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JS-10-083-357-712
     Query Match
                                     TYPE: PRT
ORGANISM: Caenorhabditis elegans
-09-903-456-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                             -09-903-456-20
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Best Local Similarity
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EQ ID NO 712
LENGTH: 82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                                                   )FTWARE: FastSEQ for Windows Version 4.0
) ID NO 20
ENGTH: 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ILE REFERENCE: 032796-090
JRRENT APPLICATION NUMBER: US/10/083,357
JRRENT FILING DATE: 2002-02-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TLE OF INVENTION: Systemic Discovery of New Genes
                                                                                                                                                                                                                                                                                                                                                                                            LICANT: Abbott Laboratories
                                                                                                                                                      OR FILING DATE: 1998-09-02
                                                                                                                                                                                                                                                                                           E REFERENCE: 6407.US.P3
                                                                                                                                                                                                                                                                                                                                                                                                                                 nt No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           083-357-712
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                                                                                                                                                                                    ENT APPLICATION NUMBER: US/09/903,456
ENT FILING DATE: 2001-07-11
R APPLICATION NUMBER: US 09/624,670
R FILING DATE: 2000-07-24
R APPLICATION NUMBER: US 09/379,095
R FILING DATE: 1999-08-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ICANT: Qiandong Zeng et al.
                                                                                                                                                                    APPLICATION NUMBER: US 09/145,828
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                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.
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INVENTION: ELONGASE GENES AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                               US20020138874AJ
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Leonard, Amanda Eun-Yeong
                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09903456
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                                                                                                                                                                                                                                                                                                                                          Huang, Yung-Sheng
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 Score 26;
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Pred. No.
DB 10;
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Length 238;
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                                                                                                                                         ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-925-301-984
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Best Local Similarity
Watches 4; Conserve
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US-10-043-487-407
                                                                      Matches
                                                                                                                                                                                           SEQ ID NO 984
LENGTH: 402
                                                                                 Query Match
Best Local Similarity
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SEQ ID NO 407
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Publication No. US20030055220A1
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                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-925-301-984
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CURRENT FILING DATE: 2002-04-30
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TLE OF INVENTION: mammalian polypeptides
116 SYTLH 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          119 SYTLH 123
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                                   1 SYTMH 5
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5. US20020052308A1
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                                                                      Conservative
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Pred. No. 1.9e
1; Mismatches
                                                                                   Score 26;
Pred. No.
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US-09-742-580-8

Sequence 8, Application US/09742580

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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Gapop 10.0 ,
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2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
n2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
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US-09-866-206A-76

US-09-956-206A-78

US-09-956-206A-78

US-09-813-398-2

US-09-795-515-11

US-09-795-515-12

US-09-795-515-14

US-09-795-515-14
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US-09-742-581-8
US-09-742-582-8
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US-10-043-487-407
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                                          Sequence 8, Appli
Sequence 712, App
Sequence 20, Appl
Sequence 407, App
Sequence 84, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 7, Appli
Sequence 77, Appli
Sequence 78, Appli
Sequence 2, Appli
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US-09-767-395-8
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US-09-880-748-1307	9-940-166A-	-09-811-384.	-09-864-761-	4-761	-866-	-09-393-	67-550-	-10-066-895-	-10-066-89	10-066-895-	066-895-	-09-956-206A-	-956-	-09-848-798-	-09-848-798-	9-795-515-3	-09-795-515-2	9-795-515-	9-795-515-2	-09-795-515-2	-09-795-515-	15-1	-09-795-515-	9-795-515-	US-09-795-515-16
1307, Ap	е : :	equence	ra -	ro -	equence	equence	equenc	ത	equence	ce 7, App	Sequence 6, Appli	63,	equence 47,	e 28	equence 27,	e 30,	e 24	equence 23,	e 22,	equence, 21,	equence 20.	equence 19	equence 18	equenc	•

ALIGNMENTS

Sequence 8, Application US/09767395
Patent No. US20020004215A1
GENERAL INFORMATION: COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/767,395
FILING DATE: 23-Jan-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/098,244
FILING DATE: <Unknown>
APPLICATION NUMBER: PCT/GB97/01835
FILING DATE: 08-JUL-1997
APPLICATION NUMBER: GB 9614292.2
FILING DATE: 08-JUL-1996
APPLICATION NUMBER: GB 9624880.2
PILING DATE: 29-NOV-1996
APPLICATION NUMBER: GB 9624880.2
PILING DATE: 29-NOV-1996 ATTORNEY/AGENT INFORMATION: NAME: David W. Clough ZIP: 60606-6402 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk CORRESPONDENCE ADDRESS APPLICANT: Osbourn, Jane K
Derbyshire, Blaine J
McCafferry, John G
Vaughan, Tristan J
Johnson, Kevin S
TITLE OF INVENTION: Labelling and selection
NUMBER OF SEQUENCES: 36 FILING DATE: 29-NOV-1996
APPLICATION NUMBER: GB 9
FILING DATE: 18-JUN-1997 CITY: Chicago STATE: Illinois REGISTRATION NUMBER: 36,107 COUNTRY: USA STREET: ADDRESSEE: Marshall, 6300 Sears 1, O'Toole, Gerstein, Murray & Borun Tower, 233 South Wacker Drive 9712818.5 Version #1.25 of molecules (EPO)

30 SYTMH

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                                                                                                                                                                                                                 O157:H7 and genomic comparison with DNA Res. 8:11-22(2001).
EMBL; AP000422; BAA94106.1; -.
EMBL; AP002554; BAB34590.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Complete nucleotide sequence of the propha
erotoxin 2 genes of the enterohemorrhagic
derived from the Sakai outbreak.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=20198780; PubMed=10734605;
Makino K., Yokoyama K., Kubota Y., Yutsudo C.H., Ki
Kurokawa K., Ishii K., Hattori M., Tatsuno I., Abe
Yamamoto K., Ohnishi M., Hayashi T., Yasunaga T., F
Sasakawa C., Shinagawa H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRAIN=0157:H7;
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EMBL; AP000363; BAA84290.1; -.
EMBL; AP004402; BAB87939.1; -.
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                                                                                                                                                          Hypothetical protein. SEQUENCE 95 AA; 10446 MW;
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SEQUENCE 95 AA; 10
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Shiga toxin 2-converting phages,";
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TRAIN=0157:H7 / R
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1-OCT-2000
                                                 Local Similarity
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SYTMH 5
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5; Conserv
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O (TrEMBLrel. 15,
2 (TrEMBLrel. 20,
al protein H0078.
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                                                                        100.0%;
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tarai M., Kobayashi M., Kano S., Hamabata T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gamma
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Pred. No.
                                                                           Score 29;
Pred. No.
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                                                                                                                                                             5F2FB7C449685BAC CRC64;
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                                                 Mismatches
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9.6;
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be H., Iida T.,
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Tobe T.,
Yasunaga T.,
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Best Local S
Matches 5
                                                                                                           InterPro; IPR001314; Chymotrypsin.
InterPro; IPR001254; Ser protease_Try.
IfterPro; IPR001254; Ser protease_Try.
Pfam; PF00089; Trypsin; I
PRINTS; PR00722; CHYMOTRYPSIN.
SMART; SM00020; Tryp SPc; 1.
PROSITE; PS001240; TRYPSIN DOM; 1.
PROSITE; PS00134; TRYPSIN HIS; UNKNOWN ]
PROSITE; PS00135; TRYPSIN_SER; 1.
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EMBL; AY060229; AAL25268.1; -.

FlyBase; FBgn0037913; CG6783.

SEQUENCE 123 AA; 14109 MW; 838C453Cpp/cmm.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; AF109648; AAD05563.1; HSSP; P20231; 1AAO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lumbricus bimastus.
Eukaryota; Metazoa; Annelida; Clitellata;
Lumbricina; Lumbricidae; Lumbricus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence up
01-JUN-2002 (TrEMBLrel. 21, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      096687
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01-MAR-2002
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Eukaryota; Metazoa; Arthropoda; Trac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (NOV-1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fibrinolytic enzyme (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GH01923p.
                                          Hydrolase; Serine protease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (NOV-1998) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bimastus.
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Siang G., Meng X., Fu
Cloning of the genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CBI_TaxID=86417;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95 SYTMH
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5; Conser
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    242 AA;
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(TrEMBLrel. 19, 1
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24837 MW;
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100.0%; Pred. No. 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S., Li L., Hou Y., Chen F., Shan F., Qin Y., encoding fibrinolytic enzymes from Lumbricus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the EMBL/GenBank/DDBJ databases TO PEPTIDASE FAMILY S1: ALSO KNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEPTIDASE FAMILY S1; ALSO KNOWN AS THE
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Last sequence update)
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                                                                                                                   UNKNOWN_1.
F912425D2724745D CRC64;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                     number of hits satisfying chosen parameters:
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Gapop 10.0 , Gapext 0.5
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1: sp_archea:*
2: sp_bacteria:*
3: sp_fungi:*
4: sp_human:*
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29
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Copyright (c) 1993 - 2003 Compugen Ltd.
sp_plant:*
sp_rodent:*
sp_virus:*
sp_vertebrate:*
sp_unclassified:*
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sp_organelle:*
sp_phage:*
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sp_mammal:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

654 221 10 9 8 7 6 5 4 2 2 1	Result No.
N N N N N N N N N N N N N N N N N N N	Score
100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 89.7 89.7 89.7 89.7 89.7	Query Match
95 242 242 388 945 945 113 113 1143 1143 158	Length 1
9 1 2 1 2 2 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	BB
Q9XJH3 Q9XXG9 Q9STC7 Q9568C7 Q8T130 Q93378 Q93189 Q9HKT2 Q	ID
Q9xjh3 bacteriopha Q9kxg9 escherichia Q95tc7 drosophila Q95tc7 drosophila Q95tc7 lumbricus b Q8ti30 methanosarc Q93378 caenorhabdi Q5223 leuconostoc Q9hkt2 thermoplasm Q33189 staphylococ Q9umi0 homo sapien Q9ug94 homo sapien	Description

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239 homo sap	8 homo sap	homo sap	8 homo sapie	3 homo sapie	homo sapien	l ralstonia	borr	μ,	caenorhabdi	ַת	4 homo sanie	0 notechis a	3 laticaud	4 homorosela	7 demansia a	pelamis pl	plethodon	plethodon w	human hero	7 ralstonia	O9b3v3 plethodon f	plethodon			Q9b3v2 plethodon a	byroc	7	O9atp4 pennisetum

ALIGNMENTS

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"Sequence analysis of Stx2-Converting phage VT2-Sa shows a great	Miyamoto H., Nakai W., Yajima N., Fujibayashi A., Higuchi T., Sato K., Matsushiro A.;	919; PubMe	SEQUENCE FROM N.A.		the EVEI (Command (Spn)	SPECIES=Phage VI2-Sa;		Submitted (FEB-1999) to the EMBL/GenBank/DDBJ databases. [3]	G. III;	Phage	SEQUENCE FROM N.A.		J. Bacteriol. 181:1767-1778 [1993]	Shiqa toxin as a phage late-came product " Becnerichia COII OI57:H7:	TYPE BEACCHER 9. X. ;	מאליב [ם ד. יד		Phage	SEQUENCE FROM N.A.		NCBI TaxID=10730, 97081 180816; will tasbitted dsbun phages.	atere include the deline	ng bacter		Bacteriophage 933W.		1010	(Tramburel, 21,	(Trembline) 12 Lagt semisors	-1999 (TrEMBLrel, 12		Q9XJH3 PRELIMINARY; PRT: 95 AA	Q9XJH3	RESULT 1

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Best Local Similarity
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SEQUENCE
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                                       Frangsmyr L. Teglund S., Israelsson A., Hammarstrom S. "Characterization of upstream promotor region, exon1 and PSG gene family.", Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases -i- SUBCELLULAR LOCATION: Secreted (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW he European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-gor send an email to license@sisb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PSG gene family."; Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases
                                                                                               SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
Teglund S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SITE
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EMBL; AF106561; AAD21024.1;
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Teglund
                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -i- SIMILARITY: CONTAINS
                                                                                                                                                                                                                                                 Jo-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Pregnancy-specific beta-1-glycoprotein 8 precursor
                                                                                                                                                                                                                                                                                                                     29UQ74;
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Mammalia; Eutheria;
                                                                                                                                                        NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                       iomo sapiens (Human)
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                                                                                                                                                                                                                                       Fragment).
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                                DEVELOPMENTAL STAGE:
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1 34 POTENTIAL.

1 34 PREGNANCY-SPECIFIC BETA-1-GLYCOPROTEIN
35 >143 PREGNANCY-SPECIFIC BETA-1-GLYCOPROTEIN
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 BELONGS
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129
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104
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Primates;
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Primates;
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                                PSBG ARE PRODUCED
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THE IMMUNOGLOBULIN SUPERFAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 26; DB 1;
Pred. No. 19;
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CELL ATTACHMENT SITE (POTENTIAL)

N-LINKED (GLCNAC. . .) (POTENTIA

N-LINKED (GLCNAC. . . ) (POTENTIA

"""""" (GLCNAC . . ) (POTENTIA
                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                  PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2C452383C8381D28 CRC64;
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                                                                                                                                                                                                                                                                                                                                  144 .AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GLCNAC. .
                               IN HIGH QUANTITY DURING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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RESULT 4
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Best Local S
Matches 4
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P27013;
01-AUG-1992 (
01-AUG-1992 (
01-NOV-1997 (
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EMBL; X65543; CAA46511.1;
PIR; S21340; S21340.
HSSP; P46088; 2GSQ.
                                                                                                                                              "Facile cloning and sequencing of S-crystallin genes from octopus lenses based on polymerase chain reaction.", Biochem. Int. 27:173-178 (1992).

-I- FUNCTION: S-CRYSTALLINS ARE STRUCTURAL COMPONENTS OF SQUIDS AND OCTOPI EYE LENS. CONTAINS RELATIVELY LITTLE GST ACTIVITY (1/1000 OF THAT OF MAMMALIAN GST ENZYME).

-I- TISSUE SPECIFICITY: LENS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institute. There are no restricted and this statement is not removed. Usage by an entities requires a license agreement (See http://www.incorsend an email to license.agreement)
                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Mollusca; Co
Incirata; Octopodidae; Octopus
NCBI_TaxID=6645;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                or send an email to license@isb-sib.ch).
                                                                       use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                    MEDLINE=92328840;
Lin C.-W., Chiou S
                                                                                                                                                                                                                                                                                                                                    Octopus vulgaris (Octopus).
Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Octopoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AF106556; AAD21022.1; EMBL; AF106555; AAD21022.1;
                                                            entities requires a license
                                                                                               the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                             TISSUE=Lens;
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(Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
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. 23, Last sequ
. 35, Last anno
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POTENTIAL
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Pred. No.
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GLCNAC. . .)
GLCNAC. . .)
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Perfect score:
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seq length: 0 seq length: 2000000000
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Match
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Gapop 10.0 ,
                         112892 seqs, 41476328 residues
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29
1 SYTMH 5
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PSG6_HUMAN
PSG6_HUMAN
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PSG1_HUMAN
PSG6_HUMAN
PSG
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     P47210 homo sapien
P11464 homo sapien
Q00888 homo sapien
Q100886 homo sapien
Q15235 homo sapien
Q15235 homo sapien
Q16557 homo sapien
Q16557 homo sapien
Q10889 homo sapien
Q00889 homo sapien
P46470 xenopus lae
P54871 caenorhabdi
Q49187 lycopersico
Q49187 lycopersico
Q82436 cucumis mal
Q98896 cucumis mal
                                                                                                                                                                                                                                                                                                                                                                             Q9uq72 homo sapien
Q9uq74 homo sapien
Q9uq74 homo sapien
P27013 octopus vul
P27011 octopus dof
Q25626 octopus dof
Q25626 octopus dof
Q2702 chlamydia p
Q92792 chlamydia p
Q92792 chlamydia m
C84538 chlamydia m
C85538 homo sapien
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6 cucumis mer
6 cucumis sat
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RESULT PSGD_H ID PSGD_H AC Q DT 3 DT 1 DT 1 DB P. GN PPG OS H	Oy Ma	SQ PT	######################################	YME AC COSS DITTOR OCCOSS DITTOR OCCOSS DITTOR	RES	
D HUMAN PSGD HUMAN PSGD HUMAN STANDARD; PRT; 143 AA. Q9UQ72; 30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update) 15-UUN-2002 (Rel. 41, Last annotation update) Pregnancy-specific beta-1-glycoprotein 13 precursor (PSBG-13) (Fragment). PSG13. Homo sapiens (Human).	Query Match 89.7%; Score 26; DB 1; Length 102; Best Local Similarity 80.0%; Pred. No. 13; Matches 4; Conservative 1; Mismatches 0; Indels 0; Gaps 0; Y 1 SYTMH 5 : 5 47 TYYMH 51	EMBL; Z48639; CAA88581.1; SGD; S0004867; YMR254C. Hypothetical protein; Transm TRANSMEM 1 21 TRANSMEM 42 62 TRANSMEM 42 62 TRANSMEM 68 88 SEQUENCE 102 AA; 11816 MW		VE Saasassassassassassassassassassassassass	35 26 89.7 3119 1 CAIC_MOUSE 36 26 89.7 3124 1 CAIC_CHICK 25 86.2 138 1 TSHB_CANFA 37 25 86.2 138 1 TSHB_HUMAN 39 25 86.2 138 1 TSHB_HUMAN 40 25 86.2 219 1 HISB_MYCSM 41 25 86.2 272 1 YEGX_ECOLI 42 25 86.2 297 1 YEGX_ECOLI 43 25 86.2 297 1 YEGX_ECOLI 44 25 86.2 297 1 YEGX_ECOLI 45 86.2 297 1 YEGX_ECOLI 46 86.2 297 1 YEGX_ECOLI 47 25 86.2 297 1 YEGX_ECOLI 48 25 86.2 297 1 YEGX_ECOLI 49 86.2 297 1 YEGX_ECOLI 40 25 86.2 297 1 YEGX_ECOLI 41 25 86.2 297 1 YEGX_ECOLI 42 25 86.2 297 1 YEGX_ECOLI 43 25 86.2 297 1 YEGX_ECOLI 44 25 86.2 297 1 YEGX_ECOLI 45 86.2 297 1 YEGX_ECOLI 46 25 86.2 309 1 YEEY_ECOLI 47 26 86.2 309 1 YEEY_ECOLI 48 27 28 86.2 309 1 YEEY_ECOLI 49 30 30 30 30 30 30 30 30 30 30 30 30 30	4 26 89.7 3063 1 CA1C_HUMAN Q99715 homo:

A;Pathway: detoxification
A;Note: this protein has weak catalytic activity
C;Superfamily: glutathione transferase
C;Keywords: detoxification; glutathione; transferase

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C;Superfamily: glutathione transferase
                                                                                                                                                                          밁
                                S-crystallin 3 - giant octopus
C;Species: Octopus dofleini (giant octopus)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 18-Jun-1999
C;Accession: C41681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3-crystallin isoform S3 -
                                                                                                                         RESULT 15
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Accession: S56762
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Date: 10-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 18-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                              Biol. Chem. 266, 24226-24231, 1991
'e: Crystallins of the octopus lens. Recruitment from detoxification enzymes. rence number: A41681; MUID:92084735; PMID:1721068
Accession: D41681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Date: 30-Jun-1992 #sequence_revision 18-Sep-1992 #text_change 18-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Species: Octopus
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Residues: 1-215 <TOM>
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Matches
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3 SYTLH 7
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.; Zinovieva, R.D.; Piatigorsky, J. 266, 24226-24231, 1991
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80.0%; Pred. No. 89;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Score 26; DB; pred. No. 89; 1; Mismatches
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A;Title: Crystallins of the octopus lens. Recruitment from detoxification enzymes.
A;Reference number: A41681; MUID:92084735; PMID:1721068
A;Recession: C41681
A;Steatus: preliminary
A;Molecule type: mRNA
A;Residues: 1-215 <TON5
A;Cross-references: GB:M65186; NID:g159732; PIDN:AAA29388.1; PID:g159733
C;Superfamily: glutathione transferase
```

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Query Match 89.7%; Score 26; DB 2; Length 215;
Best Local Similarity 80.0%; Pred. No. 89;
Matches 4; Conservative 1; Mismatches 0; Indels 0; Gaps

Qy 1 SYTMH 5
Db 3 SYTLH 7
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0

Search completed: April 16, 2003, 17:21:37 Job time: 2.54386 secs

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GenCore version 5.1.4 p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on: April 16, 2003, 17:11:36 ; Search time 1.54386 Seconds (without alignments) 311.344 Million cell updates/sec

Title: Perfect score: Sequence: US-09-644-668A-27 29 1 SYTMH 5

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

number of hits satisfying chosen parameters: 283224

283224 seqs, 96134422 residues

Minimum DB Maximum DB seq length: 0
seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 73:* 1: pir1:* 2: pir2:* pir3:*

BCOTE and is No. is the number of results predicted by chance to have greater than or equal to the score of the result being posterived by analysis of the total score distribution. being printed,

SUMMARIES

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pregnancy-specific	~		proteasome protein	ecifi	al prot	specifi	pecif	w	pregnancy-specific	specif	-spec		۲,	Ÿ	pregnancy-specific	

ALIGNMENTS

RESULT 1 G90774

hypothetical protein ECs1167 [imported] - Escherichia coli (strain O157:H7, substrair C;Species: Escherichia coli

Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 18-Jul-2001

R;Hayashi, C; Accession: G907 T., Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.

gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H. DNA Res. 8, 11-22, 2001 A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and gA;Reference number: A99629; MUID:21156231; PMID:11258796 Han,

A; Accession: G90774

A;Status: preliminary

A; Molecule type: DNA A; Residues: 1-95 < HAY>

A;Cross-references: GB:BA000007; PIDN:BAB34590.1; PID:g13360627; GSPDB:GN00154 A;Experimental source: strain O157:H7, substrain RIMD 0509952

Genetics:

8 Query Match
Best Local Similarity
Matches 5; Conserv 1 SYTMH 5 Conservative 100.0%; Score 29; DB 2; 100.0%; Pred. No. 7.7; cive 0; Mismatches 0 0; Length 95 Indels <u>,</u> Gaps 0

밁 8 SYTMH 34

RESULT 2 S17230

Ig heavy chain V region - mouse C;Species: Mus musculus (house mouse) C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 06-Jun-1997

C;Accession: S17230
R;Clackson, T.; Hoogenboom, H.R.; Griffiths, A.D.; Winter, G. Nature 352, 624-628, 1991 Nature 352, 624-628, 1991 A;Title: Making antibody fragments using phage display libraries. A;Reference number: S17230; MUID:91326098; PMID:1907718

Accession: S17230

A;Status: preliminary

A; Molecule type: nucleic acid A; Residues: 1-96 < CLA>

F;8-91/Domain:

Query Match
Best Local Similarity
Matches 5; Conserv 100.0%; ilarity 100.0%; Conservative (0; Score 29; DB 2; Pred. No. 7.8; . Mismatches 0 Length 96; Indels .. Gaps

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AAR13161

ABP38145

AAM88306

Staphylococcus epi Human immune/haema Novel human secret Pregnancy-specific

Human secreted pro N-terminal of SP1-Chlamydia pneumoni

Human secreted pro Novel human diagno C. trachomatis lip

89.7 89.7 89.7 89.7 89.7

AAY37686

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AAW05826 AAY27073

Humanised M291 ant L. bimastus plasmi Novel human diagno

Human polypeptide Human immune/haema Arabidopsis thalia

ABG18515

AAM86640 AAG59811 AA008358 AAW05825 AAW90285 AAR24107 AAW36165 AAW36167 AAY44176

Humanised M291 ant Human anti-GPIIb/I Humanised anti-Tac Humanised heavy ch Heavy chain variab MAb Fabl385 heavy

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB
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Gapop 10.0 ,
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168.037 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                               908470 seqs, 133250620 residues
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29
                                                                                                                                                                                                                                                                                                                                                                              1 SYTMH 5
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Result No.

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AAW90315 AAB67490 ABP34671 AAR21264 AAR21284 AAR21284 AAB67508 AAB67513 AAB67513

Murine Murine Light Light Heavy

Human | Human | Human | Murine

DAT: * DAT: * DAT: *	RESULT 1 AAW90315 At	IT 1 315 AAW90315 standard: Protein: 5 AA
DAT:*	AC AAW90315;	
DAT: *	DT 07-SEP-1999	(first entry)
DAT: *		****** Carc+ } /
DAT: *		Human anti-idiotypic antibody heavy chain protein CDR1 region 2.
DAT: *		
DAT: *		Antibody: GPIIb/IIIa: human: auto-antibody: anti-idiotymic: diagno
DAT: *		blood platelet membrane protein: predisposition: prevention: treat
DAT: *		autoimmune thrombocytopaenic purpura: AITP: fibrinogen binding: th
2		thrombocyte; cardiac infarction, pulmonary embolism, heavy chain;
g printed,	OS Homo sapiens	•
	PN WO9855619-A1	
	PD 10-DEC-1998.	
otion	PF 05-JUN-1998;	98WO-EP03397.
anti-idiotyp	PR 08-MAY-1998	9808+1020663
heavy chain	PR 06-JUN-1997;	97DE-1023904
ORF3644 prot		97DE-1055227
VH group 1		
VH group 1		(ASAT-) ASAT AG APPLIED SCI & TECHNOLOGY.
VH group 1	×	
chain variab		Berchtold P, Escher RFA;
chain variab		
chain variab	DR WPI; 1999-105496/09.	5496/09.
chain variab	×	

Antibody; GPIIb/IIIa; human; auto-antibody; anti-idiotypic; diagnosis; blood platelet membrane protein; predisposition; prevention; treatment; autoimmune thrombocytopaenic purpura; AITP; fibrinogen binding; thrombi; thrombocyte; cardiac infarction; pulmonary embolism; heavy chain; CDR1.

AAM93510 AAM93575 AAY57912 ABG18511 ABB58875

ALIGNMENTS

AAB68291 AAM93561

Human Human

Human

polypeptide,

Amino acid sequenc Human polypeptide, Human polypeptide,

Drosophila melanog

Caenorhabditis ele SP1-like protein e Putative polyunsat Potential polyunsa

Tomato ethylene re Tomato TGETR2 ethy

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Matches 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 80.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/01
FILING DATE: 17-JUN-1998
                                                                                                                                                                             TLE OF INVENTION
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                                                                                                                               DDRESSEE
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LICATION NUMBER: US 08/889,291
LING DATE: 08-JUL-1997
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                                                                                                                                                                                                                                                                         INFORMATION:
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PPLICATION DATA:

PPLICATION DATA:

GB 9624880.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CATION NUMBER:
. READABLE FORM:
| TYPE: Floppy disk
|:R: IBM PC compatible
                                                 60606-6402
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GY: linear
                                                                                                Chicago
                                                                                                                                                                                                                                                                                                      Application US/09375314
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                                                                               Illinois
                                                                                                             3: Marshall, O'Toole, Gerstein, Murray & Borun
6300 Sears Tower, 233 South Wacker Drive
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McCafferty, John G
                                                                                                                                                                  Johnson, Kevin S
MENTION: Labelling and Belection of molecules
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08-JUL-1997
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; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-2990
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                                                      Matches
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Best Local (
                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                       SEQ ID NO 2990
                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/064,964
                                                                                                                                                                                                                                                                                                  PPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: David W. Clough
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FILING DATE: 08-JUL-1997
APPLICATION NUMBER: PCT/
62
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                           1 SYTMH 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: GB 9712818.5
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LICATION NUMBER: GB 9614292.2
LICATION NUMBER: JUL-1996
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n Release #1.0, Version #1.25
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Pred. No.
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Pred. No.
                                                      Mismatches
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Matches
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Matches
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                                                                                                                                                                                      TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/859,583
FILING DATE: 27-MAR-1992
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NFORWATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
SECUENCES:
                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                              TELEPHONE: 415-326-2400
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                                                                                                                                                                                                                                                     (TTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 011823-004901
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                                                               Local Similarity
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                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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nes 5; Conserv
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E OF INVENTION: B-Cell Lymphoma and Cell Line
                      S HMIYS
                                                                                                                                                            amino acid
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amino acid
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                                                                                                                                                                    279 amino acida
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One Market Plaza, Steuart Tower, Suite 2000
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gingrich, Roger
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                                                                                                                                  linear
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Pred. No. 40;
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Pred. No.
                                             Mismatches
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                                                                                                                         tent No.
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NAME: David W. Clough
NUMBER OF
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APPLICATION NUMBER: GB 9614292.2

PILING DATE: 08-JUL-1996
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FILING DATE: 08-JUL-1997
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                                                                                                                                       Application US/09098244
                                       Osbourn, Jane K
Derbyshire, Elaine J
McCafferty, John G
Vaughan, Tristan J
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McCafferty, John G
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IBM PC compatible
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      Labelling and selection of molecules
                            Kevin S
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Pred. No. 20;
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                                                                                                Sequence 4, Application US/08656586
Patent No. 5834597
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                                                                                               cent No.
                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
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                                                 PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Smith, William REGISTRATION NUMBER:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                   PLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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                     LE OF INVENTION:
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LICATION NUMBER: 1
                                                                              INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                            120 amino acids
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                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                              single
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                                             , Michael S
                                                                 Yun
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                  Mutated No.
                                                                                                                                                                                                                                                  100.0%; Score 29; DB
100.0%; Pred. No. 17;
1ve 0; Mismatches
                               Claudio
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           5834597activating
 Incorporating
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Best Local Similarity 100.
Stranger 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence 8,
                                                                                                                              ZIP: 94111-3834
COMPUTER READLABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-656-586-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tent No.
FILING DATE: 31-MAY-1996
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Liebeschuetz, Joseph O.
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 11823-007210US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DAY
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Liebschuetz, Joseph O.
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 1182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
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50 SYTMH 54
                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Cole, Michael S. PPLICANT: Anasetti, Claudio
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                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                  COUNTRY:
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TOPOLOGY: linear
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Two Embarcadero Center, Eighth Floor
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Two Embarcadero Center, Eighth Floor
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100.0%; Pred. No.
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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length: 2000000000
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      262574 seqs, 29422922 residues
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29
1 SYTMH 5
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US-08-471-613-1
PCT-US93-10443-1
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US-08-714-524D-40
US-09-145-828A-13
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US-08-397-411-11
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GENERAL IN
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                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Weiner, George
APPLICANT: Gingrich, Roger
APPLICANT: Link, Brian
APPLICANT: Tso, J. Yun
                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Tso, J. Yun
IIILE OF INVENTION: Bispecific Antibody Effective
IIILE OF INVENTION: B-Cell Lymphoma and Cell Line
VUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Townsend and Townsend and Crew STREET: One Market Plaza, Steuart Tower, Suite 2000 CITY: San Francisco STATE: California
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	Sequence 13, Appl	rddw 'zr apmanhae	3 (11.	125.		14	<u>.</u>	78	Sequence 76, Appl	23	ω.	w.	37	ν	9	18,	Sequence 15, Appl	•	

ALIGNMENTS

INFORMATION:

Gingrich, Roger Link, Brian

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Query Match 100.0%; Score 29; DB 3; Length 120; Best Local Similarity 100.0%; Pred. No. 17; Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps () MULECULE TYPE: peptide US-08-397-411-10	TOPOLOGY: linear	TYPE: amino acid	 INFORMATION FOR SEQ ID NO: 10:	; TELEPHONE: 415-326-2400	; REFERENCE/DOCKET NUMBER: 011823-004901 ; TELECOMMUNICATION INFORMATION:	REGISTRATION NUMBER: 30,223	; ATTORNEY AGENT INFORMATION:	FILING DATE: 27-MAR-1992	PRIOR APPLICATION DATA:	; CLASSIFICATION: 424	FILING DATE: 01-MAR-1995	CURRENT APPLICATION DATA:	; SOFTWARE: Patentin Release #1.0, Version #1.25	; OPERATING SYSTEM: PC-DOS/MS-DOS	COMPUTER: IBM PC compatible	INTERIOR CONTRACTOR OF THE PARTY OF THE PART

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                                                                                               Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
US 08/425,336
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                                                             Local Similarity 91.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                              1 RASQSVGSSYLA 12
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RASQSVSSSYLA 35
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o. 5744580
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                                                                                                                                                                                         amino acid
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500 West Madison Street, 34th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Studnika,
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                                                                                                                                          protein
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07-JUN-1995
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19-JUN-1992
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09-DEC-1992
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12-MAY-1993
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91.7%;
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                                                                                                                                                                                                                                                                                                                 32,918
ER: 11022US07/200-70.P3.C2A
                                                             Score 49; DB 1;
Pred. No. 0.031;
0; Mismatches
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US-08-477-484B-150 Geomence 150, Application US/08477484B

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US-08-646-360-150

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Application US/08646360

Carroll, Stephen F. Studnika, Gary M.

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US-08-477-484B-150
                                                            Matches 11; Conservative
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Best Local Similarity
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                                                                                                                                                                                                TELEX: 650 388 1248
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acid
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PRIOR APPLICATION DATA:
US 08/425,336
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                        TOPOLOGY: 1
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24 RASOSVSSSYLA 35
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REFERENCE/DOCKET NUMBER: 11022US07/200-70.P3.C2A
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ZIP: 60661
                           1 RASOSVGSSYLA 12
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DATE: 18-APR-1995
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Illinois
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500 West Madison Street, 34th floor
                                                                                                                                                                                                                                                                   312/707-9155
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Carroll, Stephen F.
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUN-1995
                                                                                                                                        protein
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19-JUN-1992
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12-MAY-1993
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91.7%;
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                                                         Score 49; DB 1
Pred. No. 0.031
0; Mismatches
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Result
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Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Match
          262574 seqs, 29422922 residues
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Gapop 10.0 , Gapext 0.5
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55
1 RASQSVGSSYLA 12
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1: /cgn2_6/ptodata/1/laa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/laa/6A_COMB.pep:*

3: /cgn2_6/ptodata/1/laa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/laa/focTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/laa/backfIles1.pep:*
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                 US-08-652-558-49
US-08-4488-113B-150
US-08-448-113B-150
US-08-646-360-150
US-08-646-360-150
US-08-639-7651-42
US-08-639-7651-838-150
US-09-025-769B-37
US-08-08-053-131-183
US-08-053-131-183
US-08-053-131-183
US-08-056-762-183
US-08-086-756-147-311
US-08-862-124-17
US-08-862-124-17
US-08-899-575-147
US-08-899-575-151
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149, Appl
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160, Appl
161, Appl
183, Appl
184, Appl
17, Appl
151, Appl
151, Appl
151, Appl
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Query Match 89.1%; Score 49; DB 2; Length 107 Best Local Similarity 91.7%; Pred. No. 0.03; Matches 11; Conservative 0; Mismatches 1; Indels	CHARACTERISTICS: 107 amino acids amino acid NESS: single f: linear IYPE: protein	REGISTRATION UNDER: 30,237 REFERENCE/DOCKET NUMBER: 95,497-L TELECOMMUNICATION INFORMATION: TELEPHONE: 617-345-9110 TELEPAX: 617-345-9110 INFORMATION FOR SEQ ID NO: 49;	APPLICATION NUMBER: US/08/652,558 FILING DATE: JUNE 6, 1996 PRIOR APPLICATION DATA: APPLICATION UNMBER: PCT/IB94/00387 FILING DATE: NOVEMBER 21, 1994 ATTORNEY/AGENT INFORMATION: NAME: VANYETT LEVN B	ZIP: 02109 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch diskette, 1.4 Mb storage COMPUTER: IBM PC compatible COMPUTER: PC-DOS/MS-DOS SOFTWARE: WordPerfect 6.1 CURRENT APPLYCATION DATE: CURRENT APPLYCATION DATE:	RESSET:	RESULT 1 US-08-652-558-49 US-08-652-558-49 Sequence 49, Application US/08652558 Patent No. 5861155 GENERAL INFORMATION: APPLICANT: LIN, AUGUSTINE YEE-THARN TITLE OF INVENTION: HUMANIZED ANTIBODIES AND USES TITLE OF INVENTION: HUMANIZED ANTIBODIES AND USES	ALIGNMENTS	9 42 76.4 108 1 US-08-899-575-86 0 42 76.4 108 1 US-08-999-575-86 1 42 76.4 108 5 PCT-US95-08743-86 2 42 76.4 109 1 US-08-162-102C-23 3 42 76.4 109 5 PCT-US93-08786-23 4 2 76.4 111 1 US-08-276-852-149 5 42 76.4 111 1 US-08-899-575-149	43 /8.2 107 1 US-08-276-852-87 43 78.2 107 1 US-08-899-575-87 43 78.2 107 1 US-08-899-575-87 43 78.2 107 5 PCT-US95-08743-87 42 76.4 107 1 US-08-162-102C-38 42 76.4 105 1 US-08-276-852-93 42 76.4 105 1 US-08-276-852-93 42 76.4 105 1 US-08-899-575-93 42 76.4 105 5 PCT-US95-08743-93 42 76.4 105 1 US-08-899-575-93 42 76.4 105 1 US-08-899-575-93 42 76.4 105 1 US-08-899-575-93	8 44 80.0 129 2 US-08-480-774A-4
 2; Length 107; ; 1; Indels 0; Gaps 0		•		·Ω		AND USES	coduction 1437	Sequence 86, Sequence 86, Sequence 86, Sequence 86, Sequence 86, Sequence 23, Sequence 23, Sequence 149	Sequence 87, Sequence 87, Sequence 87, Sequence 38, Sequence 33, Sequence 93,	٠

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US-09-828-708-40
                  Query Match
Best Local Similarity
Marches 11; Conserv
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; ORGANISM: Homo sapiens
US-09-828-708-40
                                                                                                           ; ORGANISM: Homo sapiens
US-09-828-708-41
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CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 123
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 40
LENGTH: 12
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Best Local Similarity 91.7%;
Matches 11; Conservative
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ORGANISM: Homo sapiens
-09-828-708-38
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                                                                                                                                                                                                                                                                                                                                                                                                Pacent No.
                                                                                                                                                          TITLE OF INVENTION: Autoantibodies to glucose-6-phosphate isomerase and their particititle OF INVENTION: autoimmune disease
FILE REFERENCE: 1361.005US1
CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 123
SOFTWARE: FastSEQ for Windows Version 4.0
18Q ID NO 41
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ditzel, H. APPLICANT: Burton, D.
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Local Similarity. 91.7%;
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                                       Conservative
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91.7%;
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                                   Score 49; DB 10
Pred. No. 0.013;
0; Mismatches
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Pred. No. 0.013;
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; SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
; LENGTH: 74
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-125-687-23
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; ORGANISM: Homo sapiens
US-09-828-708-42
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GENERAL INFORMATION:
APPLICANT: Luo, Peter
APPLICANT: Luo, Peter
TITLE OF INVENTION: STRUCTURE-BASED CONSTRUCTION OF
FILE REFERENCE: 26050-705
CURRENT APPLICATION NUMBER: US/10/025,687
CURRENT FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 28
SOFTMARE: Patentin version 3.1
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Best Local Similarity
Matches 11; Conserv
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Publication No.
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Patent No. US20020142255A1
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APPLICANT: Luo, Pet
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Patent No. US20020146753A1
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TITLE OF INVENTION: STRUCTURE-BASED CONSTRUCTION OF HUMAN ANTIBODY LIBRARY
FILE REFERENCE: 26050-705
CURRENT APPLICATION NUMBER: US/10/125,687
CURRENT FILING DATE: 2002-04-17
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
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Local Similarity 91.7%;
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91.7%;
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                                      288829 seqs, 75613885 residues
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105
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115
246
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_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
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10 US-09-828-708-41
10 US-09-828-708-42
10 US-09-828-708-42
10 US-10-125-687-23
10 US-10-125-687-23
10 US-10-194-975-82
10 US-09-828-708-6
10 US-09-828-708-7
10 US-10-125-687-10
10 US-10-125-687-10
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256.547 Million cell updates/sec
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NEW PUB.pep: *
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                 Sequence 7,
Sequence 21,
Sequence 10,
Sequence 10,
Sequence 40,
Sequence 5,
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Sequence 82,
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Sequence 40,
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, 38	350	e 38,	Sequence 10, Appl	Sequence 92, Appl	•		5	Sequence 170 App	298	m.	776	Sequence 8, Appli	91,	14,	Sequence 905, App	e 34	1499	83	24,		88	9 1	178	4	17	Sequence 1534, Ap

ALIGNMENTS

RESULT 1

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CURRENT APPLICATION NUMBER: US/10/001,934
CURRENT FILING DATE: 2001-11-15
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 62
LENGTH: 12
TYPE: PRT
ORGANISM: Homo sapiens
US-10-001-934-62
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/ Sequence 62, Application US/10001934

/ Publication No. US20030032782A1
APPLICANT: Ditzel, H.
APPLICANT: Burton, D.
APPLICANT: Schaller, M.
APPLICANT: Schaller, M.
APPLICANT: Schaller, M.
TITLE OF INVENTION: Autoantibodies to glucose-6-phosphate isomerase and their partiparties of invention: autoimmune disease
FILE REFERENCE: 1361.005US1
CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 123
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Best Local S
Matchés 11
                                                                                                                                                                                          Sequence 38, Application US/09828708 Patent No. US20020146753A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: HUMAN PEPTIDES/PROTEINS CAUSING OR LEADING TO THE TITLE OF INVENTION: KILLING OF CELLS INCLUDING LYMPHOID TUMOR CELLS FILE REFERENCE: GPCG-P01-003
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nes 11; Conserv
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91.7%;
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                                                        Synthetic polypeptide(s) for inducing anti-idiotype antibodies useful for treating auto immune-, endocrine- and rheumatic-diseases and myasthenia gravis
                                                                                                                                                                                                                                                                                                                                                                                                       05-AUG-1985;
28-DEC-1983;
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Claim 10; Page 42;
                                                                                                                                                                                                     WPI; 1991-368612/50.
                                                                                                                                                                                                                                                                            Carson
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83US-0566172.
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44pp; English
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Pred. No.
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0.00018;
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Three rabbits were inoculated with two s.c.

injections of PGL1

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     The invention relates to a composition which includes a polypeptide or a multivalent polypeptide comprising one or more antibody-based antigen -binding domain of human composition with binding specificity for an antigen expressed on surface of a human cell, especially HLA-DR (human leukocyte antigen DR), where treating cells expressing HLA-DR with the multivalent polypeptide causes or leads to killing of cells without need of cycotoxic entities or immunological mechanisms. Also included are the nucleic acid and a host cell harbouring vector or nucleic acid. The polypeptide (optionally linked to cytotoxic or immunogenic agent), the nucleic acid and the host linked to cytotoxic or immunogenic agent), the nucleic acid and the host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sera were analysed for anti-peptide antibody activity by ELISA.
All three immunised rabbits produced anti-PGL1 antibody. Antisera
drawn from a rabbit on two different days reacted significantly with
the corresp. intact IGM-RF Glo.
See also AAR15365-74 and AAR15375.
                                                                                                                                                                                                                                                                                                                                                            Nagy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antibody-based antigen; HIA-DR; CDR; antibody; human leukocyte antigen; cell proliferative disorder; lymphoma; human leukocyte antigen disorder; lymphoma; B cell lymphoma; leukaemia; MHC class II antigen disorder; lymphoma; B cell lymphoma; leukaemia; acute myeloid leukaemia; rheumatoid arthritis; multiple sclerosis; grave's disease; insulin-dependent diabetes; transplant rejection; systemic lupus erythematosus; graft vs. host disease; myasthenia gravis; irritable bowel disease; Sjogren's syndrome; interleukin-2 secretion;
                                                                                                                                                                                                                                                       Polypeptide compositions which bind to cell surface epitopes, we multivalent form kills lymphoid tumour cells and in monovalent causes immunosuppression or inhibits activation of lymphocytes
                                                                                                                                                                                                                                                                                                                             WPI; 2002-075288/10
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06-OCT-2000; 2000US-238492P.
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OM protein - protein search, using sw model

Run on: April 16, 2003, 16:56:17; Search time 9.51579 Seconds (without alignments) 168.037 Million cell updates/sec Perfect score: 55 Sequence: 1 RASOSVGSSYLA 12

Scoring table: BLOSUM62 Gapop 10.0, Gapext 0.5

Searched: 908470 seqs, 133250620 residues

To number of hits satisfying chosen parameters: 908470 Minimum DB seq length: 0 Maximum DB seq length: 2000000000
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	Post-processing:
1: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:* 2: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:* 3: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:* 4: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:* 4: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:* 5: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:* 6: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1986.DAT:* 7: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1986.DAT:* 8: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:* 9: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:* 10: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:* 11: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:* 12: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:* 13: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1992.DAT:* 14: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:* 15: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:* 16: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:* 17: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 18: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 19: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 20: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 21: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 22: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 23: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 24: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 25: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 26: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 27: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 28: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 29: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 20: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 21: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 22: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:* 23: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:*	> E3E
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ALIGNMENTS

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Korman AJ, Halk EL, Lonberg N;	(MEDA-) MEDAREX INC.	24-AUG-1999; 99US-0150452.	24-AUG-2000; 2000WO-US23356.	01-MAR-2001.	WO200114424-A2.	Homo sapiens.	transplant rejection; graft versus host disease.	multiple sclerosis; insulin-dependent diabetes mellitus; inflammation.	rheumatoid arthritis; myasthenia gravis; lunus murron, utility;	autoimmune disease; infectious disease; inflammation, allery, cancer;	cytotoxic T lymphocyte associated antigen-4. CTLA-4. RT ligand. carrow.	Complementarity determining region. CDP, immune regnored, antibod.	Human light chain complementarity determining region 1 (CDR1).	29-MAY-2001 (first entry)	AAB67493;	T 1 493 AAB67493 standard; peptide; 12 AA.

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MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
                                                                                      Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eigimeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chilaringworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd in Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quail M.A., Rajandram M.A., Rogers J., Rutter S., Seeger K., Skelton S., Squares S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B.G.; "Deciphering the biology of Mycobacterium tuberculosis from the
                                                                                                                                                                                                                                                    STRAIN=H37RV;
                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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`1-MAR-2002 (TrEMBLrel. 20, Last an

'ypochetical protein Rv0412c.

RV0412C OR MT0425 OR MTCY22G10.08C.
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SMART; SM00406; IGv; 1.
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Mammalia; Eutheria;
 STRAIN=CDC 1551 / Fleischmann R.D.,
                                                                                                                                                                                                                               MEDLINE=98295987; PubMed=9634230;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Myosin-reactive
                                                              Complete genome sequence.";
Nature 393:537-544(1998).
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                                SEQUENCE FROM N.A.
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CBI_TaxID=1773;
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11-MAY-1997 (TrEMBLrel. 03,
11-MAY-1997 (TrEMBLrel. 03,
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OSHKOSH;
Alland D.,
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Primates;
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J.A.,
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Mycobacteriaceae; Mycobacterium
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Kolonay J.
Delcher A.
Bishai W.;
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Q9BLB1;
01-JUN-2001
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CONFLICT 355 355 D -> Y (IN REF. 2).

SEQUENCE 439 AA; 47115 MW; 7B57FA98C0472465 CRC64;
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Viruses; dsDNA viruses,
NCBI_TaxID=148603;
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Myler P.J., Sisk E., Caw
McDonagh P., Stuart K.,
Submitted (APR-2001) to
                                                            SEQUENCE FROM N.A.
STRAIN=FRIEDLIN;
                                                                                                                   Leishmania major.
Eukaryota; Euglenozoa;
NCBI_TaxID=5664;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
MEDLINE=20572070; PubMed=11123671;
MEDLINE=20572070; Jain S., Kriakov J.,
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J.F., Nelson W.C., Umayam
A., Utterback T., Weldman
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66.7%;
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Cawthra J., Handley F., V
K., Ivens A., Worthey B.A.
to the EMBL/GenBank/DDBJ
                                                                                                                                             Kinetoplastida;
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Pred. No.
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L.A., Ermolaeva M., Salzbu
J., Khouri H., Gill J., M.
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Salzberg S.
l J., Mikula
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Match
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Gapop 10.0 , Gapext 0.5
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         Q9UL86
Q9U188
P96258
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Q9BL11
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Q9A6C2
Q92VK1
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Q9ul86 homo sapien
Q9ul78 homo sapien
P96258 mycobacteri
Q9b076 mycobacteri
Q9b011 leishmania
Q9b1b1 leishmania
Q98k17 rhizobium l
Q9a6c2 caulobacter
Q985d0 rhizobium l
Q9avk1 rhizobium m
Q8y2vk1 rhizobium m
Q8y4k6 rattus norv
Q93nf6 arthrobacte
Q17844 caenorhabdi
Q8yef5 brucella me
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ALIGNMENTS

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1 RASQSVGSSYLA 12 24 RASQSVSSSYLA 35	Query Match 89.1%; Score 49; DB 4; Length 109; Best Local Similarity 91.7%; Pred. No. 0.029; Matches 11; Conservative 0; Mismatches 1; Indels 0; Gaps 0;	NON TER 109 109 SEQUENCE 109 AA; 11928 MW; 243325F72C7DAC83 CRC64;	SMART; SM00406; IGv; 1.	Pfam; PF00047; ig; 1.		P80362, 1W	Clin. Immunol. Immunopathol. 87:184-192(1998). EMBL: AF035028. BANE6264 1.	",	n-reactive autoantibodies in rheumatic carditis and		N N Down	MEDITAMENTO STATES OF THE STAT		D=9606;	Primates: Catarrhini: Hominidae:	Vortobests		active immunoglobulin ka	01-DEC-2001 (TrEMBLrel. 19, Last annotation undate)	(TrEMBLrel. 13, Creat	Q9UL86 PRELIMINARY; PRT; 109 AA. Q9UL86;	RESULT 1

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-I- DISEASE: THE PROTEIN IS ONE OF THE SURFACE IMMUNOGLOBULIN M. PATIENTS WITH CHRONIC LYMPHOCYT.
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MEDLINE=88171307; PubMed=3127527;

Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;

Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;

"Autoantibody-associated kappa light chain variable region gene
"Autoantibody-associated kappa light chain variable region gene
expressed in chronic lymphocytic leukemia with little or no some
expressed in chronic lymphocytic leukemia with little or no some
expressed in chronic lymphocytic leukemia with little or no some
expressed in chronic lymphocytic leukemia with little or no some
expressed in chronic lymphocytic leukemia with little or no some
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01-NOV-1990
01-NOV-1990
15-JUL-1999
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HSSP; P80362; 1WTL.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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HSSP; P80362; 1WTL.
                                                                                                                                                                                                       mutation. Implications for eti
J. Exp. Med. 167:840-852(1988)
-!- DISEASE: THE PROTEIN IS ON
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Mammalia; Eutheria; Primates;
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SMART; SM00406; IGv;
                                  CHAIN
                                                 SIGNAL
                                                                                SMART;
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01-NOV-1990 (Rel. 16, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-III region HIC precursor.
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SMART; SM00406; IGv; :
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91.7%;
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D IN PATIENTS WITH CHRONIC LYMPHOCYTIC
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FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
IG KAPPA CHAIN V-III REGION HIC. FRAMEWORK-1. COMPLEMENTARITY-DETERMINING-1.
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BY SIMILARITY
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COMPLEMENTARITY-DETERMINING-3.
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21-JUL-1986 (Rel. 01, Last sec
21-JUL-1996 (Rel. 38, Last and
15-JUL-1999 (Rel. 38, Last and
15-JUL-1999 (Rel. 38, Last and
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DOMAIN
              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                   DISULFID
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Pamino acid sequence of the variable regions of light chains from two range acid sequence of the variable regions of light chains from two range acids sequence of the wallow cross-reactive human IgM anti-gamma-globulins of the Wa
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                                                                 21-JUL-1986
16-OCT-2001
                                                                                      P01624;
21-JUL-1986
                                                                                                                                                                                                                                                       SEQUENCE
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SMART; SM00406; IGv; 1.
Immunoglobulin V region.
DISULFID 23 89
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PIR; A01892; K3HUSI.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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21-JUL-1986
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                                                     21-JUL-1986 (Rel. 01, Create
21-JUL-1986 (Rel. 01, Last s
16-OCT-2001 (Rel. 40, Last s
19 kappa chain V-III region
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COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
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BY SIMILARITY.
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Post-processing: Minimum Match 0%
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7E DICDI
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InterPro; IPRO0396; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
Immunoglobulin V region; Signal; Hybridoma.
NON TER 1 4
SIGNAL <1 4
CHAIN 5 100 100
DISULFID 27 93 BY SIMILARITY.
NON TER 100 100
SEQUENCE 100 AA; 10729 MW; 5D9AF363CC5263
Eukaryota, Metazoa, (
Mammalia, Eutheria, I
NCBI_TaxID=9606,
[1]
                                                                                                                                                              01-NOV-1990
01-NOV-1990
15-JUL-1999
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01-NOV-1990 (Rel. 16, Last sequence update
15-UUL-1999 (Rel. 38, Last annotation upda
Ig kappa chain V-III region HAH precursor.
Homo sapiens (Human)
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P01621;
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HSSP; P80362; 1WTL.
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Nature 307:77-80(1984).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain V-LII region NG9 precursor (Fragment).
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nilarity 91.7%;
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YV3K MUUSE
XV3K HUMAN
TRYP ASTFL
TAUC ECOLI
PROB AQUAB
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GGFI STRPY
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Pred. No. 0.0046;
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C; Accession: S34096
                                                                                                C;Accession: S67940
R;Hexham, J.M.; Furmaniak, J.; Pegg, C.; Burton, D.R.; Smith, B.R.
Autoimmunity 12, 135-141, 1992
Autoimmunity 12, 135-141, 1992
A;Tille: Cloning of a human autoimmune response: preparation and sequencing of a human A;Reference number: S67940; MUID:92314301; PMID:1617110
A;Accession: S67940
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A;Title: V-kappa gene
A;Reference number: S3
A;Accession: S34096
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ubmitted to the EMBL Data Library, September 1993; Description: Human IgM(+)IgD(+) cells, the major B
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                                         Residues: 1-91 <HEX>
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Residues: 1-91 <KLE>
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Best Local Similarity
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Residues: 1-89 <WAG>
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                      Cross-references: EMBL:X73852
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Local Similarity
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immunoglobulin V region; immunoglobulin homology
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submitted to the EMBL Data Library, K. september 1993 A;Description: Human IgM(+)IgD(+) cells, the major B A;Reference number: S37501 A;Accession: S37513 A;Status: nrelicible major basis constants.
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A;Accession: S37519
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              R;Klein, U.; Kueppers, R.; Rajewsky, K. submitted to the EMBL Data Library, September 1993 A;Description: Human IgM(+)IgD(+) cells, the major B
                                                                        Ig kappa chain V region (V-kappa 3) - human (fragment) c;Species: Homo sapiens (man) C;Jaceies: Homo sapiens (man) c;Jaceies: 06-Jan-1995 #teC;Jacession: S37517
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C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
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A;Residues: 1-92 <KLE>
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A; Residues: 1-92 < KLE>
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen_Ltd.

OM protein - protein search, using sw model

Run on: April 16, 2003, 17:11:36 ; Search time 3.70526 Seconds (without alignments) 311.344 Million cell updates/sec

Title:
Perfect score:
Sequence: US-09-644-668A-24 55 1 RASQSVGSSYLA 12

Scoring table: BLOSUM62 Gapop 10.0 ,

283224 seqs, 96134422 residues Gapext 0.5

Searched:

jumber of hits satisfying chosen parameters: 283224

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR 73:*
1: pir1:*
2: pir2:*
3: pir3:*

pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

Qy 1 RASQSVGSSYLA 12 Db 24 RASQSVSSSYLA 35	Query Match Best Local Similarity 91.7 Matches 11; Conservative	A;Residues: 1-83 «GON» C;Superfamily: immunoglobulin V region; im C;Keywords: heterotetramer; immunoglobulin	A;Reference number: A3 A;Accession: I30607 A;Status: preliminary	R;Goni, F.R.; Chen, P.P.; McGinn J. Immunol. 142, 3158-3163, 1989	C;Date: 29-Jun-1989 #sequence C;Date: 29-Jun-1989 #sequence C;Accession: I30607	RESULT 1 130607 19 kappa chain V-III r	-
	Query Match 89.1%; Score 49; DB 2; Length 83; Best Local Similarity 91.7%; Pred. No. 0.0089; Matches 11; Conservative 0; Mismatches 1; Indels 0; Gaps 0;	A;Residues: L=83 <gon> A;Residues: L=83 <gon> C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin</gon></gon>	A;Reference number: A30601; MUID:89215279; PMID:2496160 A;Recession: I30607 A;Status: preliminary A;Status: preliminary A;Status: preliminary	R;Goni, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson, D.; S. 1muunol. 142, 3188-3163, 1989	c;Abretes: nome Baptens (man) C;Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 09-May-1997 C;Accession: I30607	RESULT 1 130607 13 Kappa chain V-III region (Wei) - human (fragments)	

Ig kappa chain V region - human (fragment) C;Species: Homo sapiens (man) C;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000

R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid;
A;Reference number: S16823; MUID:91243737; PMID:1903706

A; Molecule type: mRNA A; Residues: 1-87 < BLA> A; Status: preliminary; translation not shown

A;Cross-references: EMBL:X54821
C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keyords: heterotetramer; immunoglobulin C;Keyords: immunoglobulin homology <IMM>

Matches: Query Match Best Local Similarity 1 RASOSVGSSYLA 12 Conservative 89.1**%**; 91.7**%**; 0; Mismatches Score 49; DB 2; Pred. No. 0.0094; 1; Indels

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US-08-331-398A-46
; Sequence 46, Application US/08331398A
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
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APPLICATION NUMBER: US
FILING DATE: 28-OCT-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
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RIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
PILING DATE: 30-SEP-1991
61 ADSVIGRETISRDNSKNTLYLQMNSLRAEDTAIYYCA-RTGWLGPEDYWGQGTLVTVSS, 118
                                                                                                                                                                                                                 OTHER INFORMATION:
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REGISTRATION NUMBER: 38,498
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                                                                      2VQLVBSGGGVVQPGRSLRLSCAASGFTFSSYTWHWVRQAPGKGLEWVTFISYDGNNKYY 60
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                                                                                                                                      Similarity
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FitzGerald, David
                                                                                                                                                                                                                                                                                                                                                                                          119 amino acids
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                                                                                                                86.7%; Score 549.5; DB 1;
89.1%; Pred. No. 6e-46;
Live 4; Mismatches 8;
                                                                                                                                                                                                               /note= "Human fetal immunoglobulin
56Pl'CL Variable Heavy chain (V-H)"
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Best Local Similarity
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OTHER INFORMATION:
OTHER INFORMATION:
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FILING DATE: 28-OCT-1994
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TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 015280-126120US
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                         NAME/KEY: Protein
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FILING DATE: 12-OCT-1990
                                                                                                             1 QVQLVESGGGVVQPGRSLRLSCAASGFTPSSYTMHWVRQAPGKGLEWVTFISYDGNNKYY 60
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                                                                           QVELVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY 60
ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARRSARTYYFDYWGQGTLVTVSS 119
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One Market Plaza, Steuart Street Plaza
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                     86.7%;
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Specific Antibody Fragments, Fusion Protein
                                                                                                                                                                                                                                                 /note= "Human fetal immunoglobulin
56P1'CL Variable Heavy chain (V-H)"
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                                                                                                                                                                     Score 549.5; DB 2; Length 119; Pred. No. 6e-46;
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RESULT 4 US-08-759-804A-46

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Maximum DB seq length: 200000000
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-08-331-397B-46
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Matches 108; Conservative
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                                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYTMHWVRQAPGKGLEWVTFISYDGNNKYY
                                              QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY 60
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Regult No.

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Score 558.5; DB 1 Pred. No. 8.3e-47; 2; Mismatches 6

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ZIP: 20037-3202 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy d COMPUTER: HP 9000/700 OPERATING SYSTEM: UNI: SOPTWARE: In house CURRENT APPLICATION DATA APPLICATION NUMBER: UPILING DATE: 09-SEP-1 CLASSIFICATION: 530 TELECOMMUNICATION: 520 TELECOMMUNICATION INFORM TELEPAX: (202) 293-78 TELEPAX: 6491103 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS SEQUENCE TYPE: maino acid TYPE: amino acid TYPE: amino acid TYPE: TYPE: peptide.	245-3 35, 0. 56 CANT: CANT: CANT: CANT: CANT: CANT: CANT: FR OF I OF I OF I OF I OF I OF I OF I OF I	509.5 509 508 508 508 507.5 507.5 502.5 502.5 501.5 501.5
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Best Local Similarity 86.3%;
Matches 107; Conservative
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-09-880-748-1519
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EQ ID NO 1519
                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
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TURRENT FILING DATE: 2001-06-15
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TILE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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blication No.
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olication No. US20030059937A1
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PR APPLICATION NUMBER: 60/240,816
PR FILING DATE: 2000-10-17
PR APPLICATION NUMBER: 60/276,248
PR FILING DATE: 2001-03-16
PR APPLICATION NUMBER: 60/277,379
PR FILING DATE: 2001-03-21
PR APPLICATION NUMBER: 60/293,499
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o. US20030059937A1
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Pred. No. 3.5e-35;
5; Mismatches 6
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US-09-880-748-983
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JS-09-880-748-1627
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Best Local Similarity 85.9
Matches 106; Conservative
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Best Local Similarity 84.4%;
Matches 108; Conservative
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Publication No. US20030059937A1
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FILING DATE: 2000-10-17
APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
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OF SEQ ID NOS: 3239
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Pred. No. 8.2e-35;
6; Mismatches 6
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Pred. No. 4.9e-35;
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Sequence 1627, Application US/09880748
Publication No. US20030059937A1
GENERAL INFORMATION:

PPLICANT: Ruben et al ITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

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No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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seq length: 2000000000
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Match
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634
288829 seqs, 75613885 residues
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Gapop 10.0 , Gapext 0.5
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256.547 Million cell updates/sec
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12_6/ptodata/1/pubpaa/US09 NEW PUB.pep:*
12_6/ptodata/1/pubpaa/US09 NEW PUB.pep:*
12_6/ptodata/1/pubpaa/US10 NEW PUB.pep:*
12_6/ptodata/1/pubpaa/US10 PUBCOMB.pep:*
12_6/ptodata/1/pubpaa/US10 PUBCOMB.pep:*
12_6/ptodata/1/pubpaa/US60 NEW PUB.pep:*
13_6/ptodata/1/pubpaa/US60 PUBCOMB.pep:*
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6_NEW_PUB.pep:*
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                                924, App
983, App
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1201 Ap
1627 Ap
1109 Ap
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2055, App
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977, App
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ALIGNMENTS

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; ORGANISM: Homo sapiens
US-09-880-748-924
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PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
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US-09-880-748-924
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SEQ ID NO 924
LENGTH: 247
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
PRIOR FILING DATE: 2000-06-15
115 TVSS 118
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                                                                                                                          QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
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                                                ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARPSYDILTGYLYYFDYWGGGTLV
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0. US20030059937A1
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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168.037 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                             gcgdata/geneseq/genese
                                                                                                                                                                                                                                                                                                                                                        geneseq
                                                                                                                                                                                                                                                                                               /geneseqp-embl/AA1990.DAT:*
/geneseqp-embl/AA1991.DAT:*
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-embl/AA1986.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

10 8	ገዕጠቅጠንኮ	Result No.
566.5 564	566.5 5634	Score
89.4 89.4	100.0 100.0 98.3 98.3 89.4 89.4	Query Match I
4 2 2 5 5 3 5 5 3 5	118 118 118 118 118 247 225	Length DB
222		:
AAB75002 AAB75013 AAY93701	AAB67508 AAB67513 AAB67509 AAB67514 ABB44913 AAB99380	ID.
Anti-IL® monoclona Anti-IL® monoclona Anti-IL® monoclona The heavy chain of	Light chain variab Heavy chain variab Light chain variab Light chain variab Heavy chain variab Human BLyS binding Human interleukin Human interleukin	Description

545	546	546.5	547	547.5	549	549	549.5	550	51	552.5	5	52	ŭ	554	554	554	54	4	55	tп	557.5	558	558	32	558.5	ž	o,	60.	•	61.	13	ð	564	
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223	122	248	137	119	249	118	119	252	115	225	225	225	225	490	254	118	225	225	519	252	119	463	463	225	225	120	254	225	225	225	225	252	463	463
20	22	23	23	22	23	23	17	23	21	22	22	22	22	21	21	۲2 پ	22	22	23	23	22	21	21	22	22	15	23	22	22	22	22	23	21	21
AAY08598	AAB72883	ABP45410	ABB77063	AAB67516	ABP45098	ABG30463	AAR95216	ABP45616	AAB39519	AAB75018	AAB75014	AAB99385	AAB99381	AAY56637	AAY56287	ABB07230	AAB75022	AAB99389	AAU81993	ABP45190	AAU02515	AAY93732	AAY93707	AAB75000	AAB99367	AAR52064	ABP44972	AAB75020	9938	7502	B9938	4550	Y937	AAY93727
human	anti-	BLyS bindi	ACZ885 antibody he	chain	BLyS bind		foeta	n BLyS	IL12 a	Anti-IL8 monoclona	Ħ		interleu	σ	HCAT1 clone 25 ant	끊	-		Human secreted pro	Human BLyS binding	Anti-adipocyte mon	The heavy chain of	eav	н		Heavy chain variab		н	an	Anti-IL8 monoclona	inter	an BLyS bindin	5	The heavy chain of

3310987832

ALIGNMENTS

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	AAB67508	7508	RESULT 1	•
	standard;			
	peptide;			
	118			
	\$			

Light chain variable region of anti-CTLA-4 antibody 10D1.

29-MAY-2001 AAB67508;

(first entry)

Complementarity determining region; CDR; immune response; antibody; cytotoxic T lymphocyte associated antigen-4; CTLA-4; B7 ligand; cancer autoimmune disease; infectious disease; inflammation; allergy; rheumatoid arthritis; myasthenia gravis; lupus erythematosus; multiple sclerosis; insulin-dependent diabetes mellitus; inflammation; transplant rejection; graft versus host disease. ligand; cancer;

Homo sapiens.

Ş	Ţ	Ϋ́	H	Ŧ	Ħ	Ŧ	꿒	Š
		Region		Region		Region	Key	
	/note= "CDR3"	99107	/note= "CDR2"	5066	/note= "CDR1"	3135	Location/Qualifiers	

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01-MAR-2001.

X K X B X R X 24-AUG-2000; 2000WO-US23356.

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Best Local S
Matches 105
                                            (1)
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     crausberg R.;
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases
EMBL; BC020240; AAH20240.1; -
InterPro; IPR003599; Ig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF00047; ig; 5.

SMART; SM00409; IG; 2.

SMART; SM004007; IGC1; 4.

SMART; SM00406; IGV; 1.

PROSITE; PS00290; IG_MHC; UNKNOWN_3.
                                                                                                                                                                                                                                                           Q9UL90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00290; IG
Hypothetical protein
SEQUENCE 613 AA;
                                                                                                                                    Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                       NCBI_TaxID=9606;
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                           Myosin-reactive autoantibodies in rheumatic carditis
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                                                                                                                                                                                                                                                                                                                                                                                    118
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                                                                                                                                                                                  Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DSVKGRFTISRDNSKNTLYLOMNSLRAEDTAIYYCARTGWLGPFDYWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                   S 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         105;
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IPR003006; Ig_MHC.
IPR003596; Ig_v.
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(TremBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                           PRELIMINARY;
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Primates;
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Pred. No. 1
87:184-192 (1998)
                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; V
Catarrhini;
                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60C7F5950671E315 CRC64;
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                                                         Kalis N.N.,
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                                                         Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 613
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RESULT 09UL84

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Q9UL84

PRELIMINARY;

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Best: Local Similarity
Matches 102; Conserv
                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 95; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP00047; ig; 1.
SMART; SM00406; IGv; 1.
NON TER 1 1
SEQUENCE 147 AA; 15768 MW;
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NON TER
SEQUENCE
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MEDLINE=96071149; PubMed=7475288;
Cao J., Vescio R.A., Rettig M.B., Hong C.H.,
Lichtenstein A.K., Berenson J.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR003006; Ig_MHC
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
MCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; S80860; AAD14339.1; -. HSSP; P01772; 2FB4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myeloma using PCR with patient-specific
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121
                                             113 LVTVSS 118
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                                                                                              AGSVKGRFTISRDNSKNTLYLOMTSLRVEDTAVYYCAKDGNYFDSVGYYYAGIDYWGQGT
                                                                                                                                              ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAIYYCARTG------WLGPFDYWGQGT 112
                                                                                                                                                                                                 QVHLVESGGGVVQPGKSLRLSCEASGFTFSTYGMSWVRQAPGKGLDWVALISYDGSTQYY
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113 AA;
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(TrEMBLrel.
(TrEMBLrel.
(Fragment).
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Last annotation update)
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Pred. No. 1.5e
5; Mismatches
                                                                                                                                                                                                                                                                                                  Score 497; DB 4;
Pred. No. 1.2e-42;
B; Mismatches 15
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                          8489FCAAA7BC925C CRC64
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                                                                                                                                                                                                                                                                                                       15;
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Regult
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Sequence:
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                                                                                                          481.5
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Gapop 10.0 , Gapext 0.5
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634
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sp_vertebrate:*
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Q9u1871 homo
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Q96k68 h
Q920e7
Q99ka4
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MEDLINE=77070267; PubMed=1002129;
Dreker L., Schwarz J., Reichel W., Hilschmann N.;
"Rule of antibody structure. The primary structure of a monoclonal IgG1 immunoglobulin (myeloma protein Nie), I: Purification and characterization of the protein, the L- and H-chains, the cyanogen bromide cleavage products, and the disulfide bridges.", Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540(1976).

-!- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM AN IGG1 MYELOMA
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
"mmunoglobulin V region.
                                                                                                                                                                                                          "The rule of antibody structure. The primary structure of a monoclonal IgG1 immunoglobulin (myeloma protein Nie). III. The chymotryptic peptides of the H-chain, alignment of the tryptic peptides and discussion of the complete structure."; Hoppe-Seyler's Z. Physiol. Chem. 357:1571-1604(1976).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini;
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1-JUL-1986 (Rel.
5-JUL-1999 (Rel.
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lomo sapiens (Human
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The switch point in mu heavy chains of human IgM immunoglobulins.";
Biochemistry 13:2482-2498(1974).
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73.0%;
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Pred. No. 3.6e-43;
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PIR; A02054; G1HUHL.

HSSP; P01772; 2FB4.

InterPro; IPR003006; 1

InterPro; IPR003596; 1
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P01771;
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MEDLINE=79124695; PubMed=420800;
MEDLINE=79124695; PubMed=420800;
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Mammalia, Eutheria; Primates; Catarrhini; Hominidae; Homo.
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SMART; SM00406; IGv; 1.
Immunoglobulin V region.
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21-JUL-1986
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SMART; SM00406; IGv; 1.
Immunoglobulin V region.
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Title: Perfect score: Sequence: Regult No. Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Database : Minimum DB seq length: 0
Maximum DB seq length: 200000000 Scoring table: Run on: OM protein number of hits satisfying chosen parameters: Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. 408.5 405.5 403.5 400.5 399.5 413 412.5 411 414 413.5 423.5 421 419.5 417.5 501 497.5 486.5 486 protein search, using sw model BLOSUM62 Gapop 10.0 , Gapext 0. US-09-644-668A-17 634 $\begin{array}{c} \mathbf{7} \\ \mathbf{83} \\ \mathbf{93} \\$ Match 1 QVQLVESGGGVVQPGRSLRL.....TGWLGPFDYWGQGTLVTVSS 118 April 16, 2003, 16:57:42; Search time 23:186 Seconds (without alignments) 211.085 Million cell updates/sec Query 112892 segs, 41476328 residues SwissProt_40:* GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd. Length DB HV31_HUMAN
HV36_HUMAN
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RESULT 2
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AC P01769;
DT 21-JUL-1986
DT 21-JUL-1986
DT 15-JUL-1999

STANDARD;

122 AA

(Rel. 01, Created)
(Rel. 01, Last sequence update)
(Rel. 38, Last annotation update)

Similarity 79.5%; Pred. No. 1.2e-45; 97; Conservative 12; Mismatches 9; OLVESGGGVVOPGRSLRLSCAASGFTFSSYTMHWVROAPGN:	ENCE. IMS=81013859; PubMed=6774332; IMS=81013859; PubMed=6774332; an D.W., Putnam F.W.; an D.W., Putnam F.W.; an D.W., Putnam F.W.; an D.W., Putnam F.W.; no acid sequence of the variable region of a human mu tion of a possible JH segment."; .Natl. Acad. Sci. U.S.A. 77:3229-3243(1980). INSCELLANEOUS: THIS MU CHAIN WAS ISOLATED FROM THE PLI PATIENT WITH MACROGLOBULINEMIA. A02051; M3HUAM. A02051; M3HUAM. A02051; M3HUAM. A02051; M3HUAM. A02051; M3HUAM. A02051; IPR003006; Ig_W. PPTO; IPR003006; Ig_W. PPTO; IPR003006; Ig_W. PPTO; IPR003596; Ig_V. PPTO; IPR0047; ig; 1. PPTO; SM00406; IGV; 1. POSSIBLE TO STATE TO	HUMAN HV3G HUMAN STANDARD; PRT; 122 AA. P01768; 21-JUL-1986 (Rel. 01, Created) 21-JUL-1986 (Rel. 38, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) If heavy chain V-III region CAM. Homo sapiens (Human). Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; NCBI_TaxID=9606;	34 397 62.6 111 1 HV35 MOUSE PO18 35 396.5 62.5 113 1 HV34 MOUSE PO18 36 396 62.5 122 1 HV20 MOUSE PO17 37 393.5 62.1 113 1 HV29 MOUSE PO17 38 393.5 62.1 116 1 HV39 MOUSE PO17 39 393.5 62.1 117 1 HV17 MOUSE PO17 40 392.5 61.9 117 1 HV41 MOUSE PO17 41 392.5 61.8 117 1 HV55 MOUSE PO18 42 390.5 61.6 123 1 HV25 MOUSE PO18 43 390 61.5 118 1 HV3V HUMAN PO17 44 389.5 61.4 113 1 HV39 HUMAN PO17 45 389 61.4 119 1 HV39 HUMAN PO17
Indels 4; Gaps 1; GLEWYTFISYDGNNKYY 60 : : GLEWVAVISYBGBEXYY 60 GMLGPFDYWGOGTLYTV 116 GMLGPFDYWGQGTLYTV 120 GBYRAFNYWGQGTLYTV 120	man mu THE PLA ACID. ACID.	" ы	P01804 mus musculu P01803 mus musculu P01789 mus musculu P01798 mus musculu P01798 mus musculu P017186 mus musculu P01811 mus musculu P18526 mus musculu P18526 mus musculu P01794 mus musculu P01793 mus musculu

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A;Map position: 14932.33-14932.33
C;Superfamily: immunoglobulin V region; immu C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IM
                                                           A;Status: preliminary; nucleic
A;Molecule type: mRNA
A;Residues: 1-118 <RAA>
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R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau,
R;Cuisinier, A.M.; Gauthier, J.; Boubli, L.; Fougereau,
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                                                                                                                                                                                                                          Species: Homo sapiens (man)
Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
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;Note: the nucleotide sequence was submitted to the EMBL Data Library, Superfamily: immunoglobulin V region, immunoglobulin homology
                                               Cross-references: EMBL:X62966
                                                                                                                                                                                       Accession: S31116
Kaaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Species: Homo sapiens (man)
Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
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Residues: 1-135 <CUI>
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Pred. No. 3.
                                                                                                   sequence not shown;
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A;Molecule type: mRNA
A;Residues: 1-122 <RNA>
A;Cross-references: EMBL:X62967
A;Cross-references: EMBL:X62967
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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A;Title: Restricted utilization of germ-line V(H)3 genes A;Reference number: S31104; MUID:92111633; PMID:1730252 A;Accession: S31117
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R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Esubmitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human
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R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol,
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C;Sate: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
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C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change
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                     87.1%;
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  Score 552; DB 2
Pred. No. 2e-42;
3; Mismatches
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Pred. No. 1.8e-42;
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Pred. No. 1.3e-42;
                                                                                                                                                                                                                           sequence
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  8
                                        Length 122;
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QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYTMHWVRQAPGKGLEWVTFISYDGNNKYY 60

*	GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen_Ltd.	٠.
OM protein - pro	OM protein - protein search, using sw model	
Run on:	April 16, 2003, 17:11:36 ; Search time 36.4351 Seconds (without alignments) 311.344 Million cell updates/sec	
core:	US-09-644-668A-17	
•	TO THE PROPERTY OF THE PROPERT	
Scoring table:	BLOSUM62 Gapop 10.0 , Gapext 0.5	
Searched:	283224 segs, 96134422 residues	-
To number of	number of hits satisfying chosen parameters: 283224	
Minimum DB seq l Maximum DB seq l	Minimum DB seq length: 0 Maximum DB seq length: 2000000000	
Post-processing:	Post-processing: Minimum Match 0%	

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Database :

Maximum Match 100% Listing first 45 summaries

PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

SUMMARIES

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111	120	109	119	114	123	151	133	109	128	122	130	114	130	139	133	119	132	120	122	114	134	140	122	137	118	135	121	121	Length	
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78.5	78.6	78.6	78.7	79.0	79.0	. 79.3	79.4	79.6	79.7	79.8	79.9	80.0	80.2	80.2	80.3	
124	119	113	108	136	122	119	140	117	118	120	123	143	119	111	122	
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S20782	D36005	S38490	PH1642	S31587	M3HUGA	S31107	S31686	S78486	S31677	S48798	S31114	S23624	S31108	PH1643	869910	
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ALIGNMENTS

PCCVAPARACCOCIEGRA B Q B Q BCCCAPARACCOCIIS		•	
Deavy Chain V region (VH3DJH4) - human	chain V region (M74) - human 8: Homo sapiens (man) 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change ion: G36005 der Jr., H.W.; Wang, J.Y. tl. Acad. Sci. U.S.A. 87, 6146-6150, 1990 Preferential utilization of conserved immunoglobulin he nce number: A36005; MUID:90349571; PMID:2117273 ion: G36005 i. preliminary le type: mRNA es: 1-121 <sch> references: GB:M34031 cs: GDB:IGH@; IGHDY1</sch>	Duery Match B9.4*; Score 566.5; DB 2; Length 121; Best Local Similarity 89.3*; Pred. No. 9.9e-44; Matches 108; Conservative 5; Mismatches 5; Indels 3; Gaps 1 OVOLVESGGGVVOPGRSLRLSCAASGFTESSYTMHWVRQAPGKGLEWVTFISYDGNNKYY 60	n V region (VH3DJH4) - human mo sapiens (man) n-1993 #sequence_revision 22-Jan-1993 #text S19666 ; Hoogenboom, H.R.; Bonnert, T.P.; McCaffer 222, 581-597, 1991 assing immunization. Human antibodies from umber: S19663; MUID:92085276; PMID:1748994 ps: mRMR -121 <marp <imm="" embl:x61646;="" ences:="" eterotetramer;="" homology="" immunoglobulin="" n:="" nid:937680;="" nid:937688;="" pidn:caa438=""></marp>

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US-09-240-274-178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 2
US-09-240-274-178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 42, Application US/08232081B Patent No. 5886152
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SEQ ID NO 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 178, Application US/09240274 Patent No. 6255455
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NUMBER OF SEQ ID NOS: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENT APPLICATION NUMBER: US/09/240,274
LENT FILING DATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
EARLIER APPLICATION NUMBER: 60/028,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FTIE REFERENCE: 09596-42U2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                          ZIP: 22040-0747
COMPUTER READRALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: NAKATANI, TOMOYUKI
APPLICANT: GOMI, HIDGYUKI
APPLICANT: WIJDENES, JOHN
APPLICANT: NOGUCHI, HIROSHI
                                                                                                                                                                                                                                                                                                                                      ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64 SGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 108
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                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                         TLE OF INVENTION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 107
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                                                                                                                                                                                                                                                           PALLS CHURCH
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NOGUCHI, HIROSHI
NVENTION: HUMANIZED B-B10
                                                                                                                                                                                                           USA
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                        PatentIn Release #1.0, Version #1.30 ICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95.1%; Score 538; DB 4; Length 108; 97.1%; Pred. No. 1.8e-42;
US/08/232,081B
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Matches
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                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/425,336
FILING DATE: 18-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 20 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,113B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Carroll, Stephen F. APPLICANT: Studnika, Gary M.
                                                                                            APPLICATION NUMBER: US 0 FILING DATE: 09-DEC-1992
                                                                                                                                                                                                                                                 FILING DATE: 07
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (703)
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CLASSIFICATION:
                                             APPLICATION NUMBER: US 07/901;707 FILING DATE: 19-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 94.7%;
Local Similarity 96.3%;
les 104; Conservative
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5. 5744580
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                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.25
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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MBER: US 07/787,567
04-NOV-1991
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                                                                                                              US 07/988,430
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Pred. No. 2.7e-42;
1; Mismatches 3
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Gapop 10.0 , Gapext 0.5
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Listing first 45 summaries
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/cgm2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgm2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgm2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgm2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgm2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
                                                              US-08-082-842A-14
US-07-634-278-86
US-08-477-728-86
US-08-474-040-86
US-08-484-537-86
US-08-8-99-575-86
US-08-899-575-86
US-08-862-124-17
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-477-484B-150
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                                                                                                                                                                                                                                                                             Sequence
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e 16, Appl

e 47.8, Appl

e 150, Appl

e 14, Appl

e 16, Appl

e 17, Appl

e 19, Appl

e 19, Appl

e 19, Appl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ge, Liming
APPLICANT: Moroney, Simon
APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT:
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487	487	487	487	487	489.5	489.5	489.5	489.5	491	491	492.5	493	493	501	502	502	502
86.0	86.0		96.0	86.0	86.5	86.5	86.5	86.5	86.7	86.7	87.0	87.1	87.1	88.5	88.7	88.7	88.7
107	104	104	104	104	109	109	109	.109	109	109	134	110	110	129	108	108	108
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US-08-276-852-90	PCT-US95-08743-100	US-08-899-575-100	US-08-899-575-100	US-08-276-852-100	PCT-US95-08743-147	US-08-899-575-147	US-08-899-575-147	US-08-276-852-147	PCT-US93-08786-23	US-08-162-102C-23	US-08-405-034-4	US-09-025-769B-47	US-09-025-769B-30	US-08-480-774A-4	PCT-US95-08743-99	US-08-899-575-99	US-08-899-575-99
	5	100	10.	1	Sequence 147. App	4,	14.	14	2	2	4	47	Sequence 30. Appl	4 ,	9	Sequence 99. Appl	Sequence 99. April

Pack, Peter Ilag, Vic Knappik,

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; MOLECULE TYPE:
US-09-025-769B-16
Query Match
Best Local Similarity
Matches 104; Conserv
                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                TYPE: aming
STRANDEDNESS:
Inear
                                                                                                                                                                                                                                                                                                                 FILING DATE: 18-FEB-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: MO
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                    FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                          NAME: James F. Haley, REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                               APPLICATION NUMBER: EP 9:
                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                    109 amino acids
                                                                                                                                                                            (212)596-9090
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                                                                        protein
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            95.4%;
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           Score 540; DB 4;
Pred. No. 1.2e-42;
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                       Length 109;
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Mismatches

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Best Local Similarity 93.5
Matches 101; Conservative
                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0
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PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR PELICATION NUMBER: 60/276,248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  quence 1264, Application US/09880748 blication No. US20030059937A1
                                                                                                                                                ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ILE REFERENCE: PF523
                                                                                                                                                                                                       FTWARE: Pate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E REFERENCE: PF523
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RENT APPLICATION NUMBER: US/09/880,748
RENT FILING DATE: 2001-06-15
OR APPLICATION NUMBER: 60/212,210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 ETTLTQSPGTLSLSPGERATLSCRASQSVRNSYLAWYQQKPGQAPRLLIYGASSRATGIP 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             205 DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPVTFGGGTKVEIK 252
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                                                               Local Similarity
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1 BIVITOSPGTLSLSPGERATLSCRASQSVGSSYLAWYQQKPGQAPRLLIYGAFSRATGIP 60
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/240,816 FILING DATE: 2000-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 2001-03-16
APPLICATION NUMBER: 60/277,379
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/293,499
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                                                                                                                                                                                                                                                                                                                                                                               PPLICATION NUMBER: 60/276,248
                                                                                                                                                                                                                                                                                                                                                           LING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                 LING DATE:
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                                                                                                                                                                                                                                     PatentIn
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Pred. No. 8.4e
1; Mismatches
                                                             Score 514.5;
Pred. No. 1.2
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                                         Indels
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Patent No. US20020016445A1
GENERAL INFORMATION:
                                                                                                  Sequence 5, Application US/09782397 Publication No. US20030021779A1
                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 19-APR-1996
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 17
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                      64
                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: gir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/635,109 FILING DATE: 19-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                                                                                                                                                                                                                                                                                      4 LTQSPGTLSLSPGERATLSCRASQSVGSSYLAWYQQKPGQAPRLLIYGAFSRATGIPDRF 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LE OF INVENTION:
                                                                                                                                                                                                     SGSGSGTDFTLTISRLEPEDFAVYYCQLYGNSRWTFGQGTKVEIK 107
                                                                                                                                                                                                                                      SGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 108
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                                                                                                                                                                                                                                                                                                                                         98;
                                                                               INFORMATION:
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90 MIDDLEFIELD ROAD, SUITE 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         107 amino acids
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ALLANDER, TOBIAS E
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                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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93.3%;
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HUMAN MONOCLONAL ANTIBODIES SPECIFIC
HEPATITIS C VIRUS (HCV) E2 ANTIGEN
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Pred. No. 6.8e-30;
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NUCLEOTIDES ENCODING
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                 288829 seqs,
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566
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(without alignments)
256.547 Million cell updates/sec
                                                                                                                                          Published_
                                                                                                                                                                                 Maximum Match 100%
Listing first 45 summaries
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PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.

Query Score Match Length DB

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Sequence 10, App	Sequence 10. Appl	dw 'eser apmanhae		Sequence 38. App	ddw 'orr antenna	1	Sequence 40. Appl	sequence /, Appl	ddw 're gamanhao	Semience 34 Ann	Segmence 10 April	Sequence 1219, Ap	rddw 'c aprrenhae	Total Constitution		Sequence 14, Appl	Tddv ', T apriented	110000000000000000000000000000000000000	Λ.	Sequence 8. Appli	Sequence 1264, Ap	Tw 'ccar animahan		Semience 178 Ann		Description

US-09-880-748-1499; Sequence 1499; Application US/09880748; Publication No. US20030059937A1

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                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: anti-Rh(D) antibody clone SH52
US+09-848-798-178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-848-798-178
                                                                                                                                                                                        Matches 102;
                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. SEQ ID NO 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2001-05-04
PRIOR ADPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/848,798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Siegel, Donald L.
                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
FIRE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
           63
                                                                                                                   4 LTQSPGTLSLSPGERATLSCRASQSVGSSYLAWYQQXDGQAPRLLIYGAFSRATGIPDRF 63
                                 SGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 108
SGSGSGTDFTLTTSRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 107
                                                                                       LTOSPGTLSLSPGERATLSCRASOSISSSYLAWYOOKPGQAPRLLIYGASSRATGIPDRF 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09848798
                                                                                                                                                                                        Conservative
                                                                                                                                                                                                         95.1%;
97.1%;
                                                                                                                                                                                                         Score 538; DB 9; Length 108; Pred. No. 1.4e-31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences for the variable regions of human monoclonal antibodies which are immunoreactive with human immunodeficiency virus-1 (HIV-1) envelope glycoprotein gp120. These can be used in diagnosis and therapy of HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus-1; HIV-1; human monoclonal antibody; envelope glycoprotein; gp120; diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human monoclonal antibody immunoreactive with human immunodeficiency virus-1 glycoprotein gp120, useful for detecting HIV-1 in biological sample and providing passive immunotherapy to HIV-1
                                                                                                                                                                                                                                         Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; tomato; monkey; dog; sea urchin; expressed sequence tag; EST diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                              Human EST
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25-JAN-2600; 2000US-0491404
17-JUL-2000; 2000US-0617746
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                                                                                                                                                                                                     Homo sapiens
                                                                 25-JAN-2001; 2001WO-US02687
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                                                                                                                                                                                                                                                                                                                                                            encoded protein SEQ ID NO: 1626.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein;
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Pred. No. 1.7e-33;
1; Mismatches 3
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RESULT 7
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SXXX
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Best Local Similarity
Matches 105; Conserv
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10-APR-1998;
27-JUN-1997;
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N-PSDB; AAH98760.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               urchin and tomato. These were derived from expressed sequence the from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated polypeptide for treatment of diseases, antibodies and research use -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tang YT, Liu C, Zl
Cao Y, Drmanac RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-AUG-2000; 2000US-0631451.
15-SEP-2000; 2000US-0663870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 20; Page 1102-1103; 1275pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                     Human; anti-Rh(D) a
red blood cell; Rh
                                                                                                                                                                                                                                                                                                                                             Human anti-Rh(D)
                                                                                                                                                                                                                                                                                                                                                                         14-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                               AAG93666 standard; Protein; 108 AA
                                                                                                                                                                                                                                                                        Homo sapiens
                                                                   Siegel
                                                                                                                                                                                    29-JAN-1999;
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                                                                                                  PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                            96US-0028550.
98US-0081380.
97US-0884045.
                                                                                                                                                                                    99US-0240274
                                                                                                                                                                                                                                                                                                                                             antibody clone SH52 protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhou P, Qian XB,
A, Zhang J, Werh
                                                                                                                                                                                                                                                                                                     antibody; Rh(D) binding protein; immunostimulant;
h phenotype; diagnosis; therapeutic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95.2%;
97.2%;
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Pred. No. 5.1e-33;
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Werhman T;
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New isolated protein, WPI; 2001-388931/41. N-PSDB; AAH68723.

preferably a human

anti-Rh(D) antibody for use in

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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on: April 16, 2003, 16:56:17 ;—Search time 85.6421 Seconds (without alignments) 168.037 Million cell updates/sec

Perfect score: Sequence: US-09-644-668A-7 566

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5 1 EIVLTQSPGTLSLSPGERAT......CQQYGSSPWTFGQGTKVEIK 108

number of hits satisfying chosen parameters:

908470

908470 seqs, 133250620 residues

Searched:

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database:

A_Geneseq_101002:* .: /SIDS2/gcgdata/ /SIDS2/gcgdata/geneseq/geneseqr-emb1/AA198.DAT: *
/SIDS2/gcgdata/geneseq/geneseqr-emb1/AA198.DAT: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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536 536	538 537.5	539	540	541	541.5 541	Score
94.7 94.7	95.1 95.0	95.2 95.2	95.4	95.6	95.7 95.6	Query Match Length DB
384 384	108	108 384	129	235	109	ength I
22 22	22	222	14	21	23	₩
AAU14461 AAU14462	AAG93666 ABB07231	AAB62770 AAM24101	AAR38672	AAY93729	ABB07233	ij
Human novel protei	Human anti-Rh(D) a Anti-IL-4 receptor	Human HIV-1 monocl Human BST encoded	vk325-Jk2. Homo s	The kappa chain of	Anti-IL-4 and IL-1	Description

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AAIY8226	AAY95117	AAW01265	AAR54307	AAB62768	AAW31722	ABP45253	AAB75009	AAB99376	ABG30495	ABG30492	AAB69689	AAW58493	AAR38593	AAB75023 .	AAB99390	AAU14227	AAU14225	ABP45488	AAB75024	AAB99391	AAB75011	AAB99378	AAB75027	AAB99394	AA021549	ABB07229	AAW84096	AAB75030	AAB75007	AAB99397	AAB99374	AAB62773	AAU14464	AAU14463	
Anti-gp120 antibod	Anti-gpl20 antibod	VL region of HIV n	Anti-HIV gp120 imm	Human HIV-1 monocl	Alpha light chain	Human BLyS binding	н	interleuk		Human anti-CD40 mo	Humanised CMV5 ant	Human kappa light	Human lambda light	Anti-IL8 monoclona	interleukin	Human novel protei	Human novel protei	Human BLyS binding	Anti-IL8 monoclona	Human interleukin	Anti-IL8 monoclona	Human interleukin	Anti-IL8 monoclona	Human interleukin	Antibody screening		< ×	8TI	н	inter	interl	HIV-1 monoc	Ë	Human novel protei	-

ALIGNMENTS

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D ABB07233 st	7233	LT 1
standard;		
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otein; 109 A

ABB07233;

26-MAR-2002 (first entry)

Anti-IL-4 and IL-13 receptors MAb 27A1 light chain variable region.

Human; antibody; interleukin; IL-4; antagonist; receptor; IL-4 receptor; antiarthritic; dermatological; antiulcer; antiinflammatory; cytostatic; antisickling; immunosuppressive; tuberculostatic; ophthalmological; IL-13; antianemic; antithyroid.

Homo sapiens.

PR	g X	\$ B \$	X P	E X	F.	P P	FŢ	ΤΉ	FH	Ķ
26-MAY-2000; 2000US-0579808 19-SEP-2000; 2000US-0665343	25-MAY-2001; 2001WO-US17094.	06-DEC-2001.	WO200192340-A2.		Region	Region		Region	Key	
00US-057	CSD-OMIO			/note=	9099	5157	/note=	2435	Locatio	
79808. 55343.	17094.			<pre>/note= "complementarity determining region (CDR)</pre>	Compacinoment at y	700tes "Complementarity determining region	<pre>/note= "complementarity determining region (CDR)</pre>		Location/Qualifiers	
				determining	A COLUMN	determining	determining			
				region		100	region			
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13

DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 108

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Best Local S
Matches 99
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                       Clin. Immunol. Immunopathol. EMBL; AF635031; AAD56267.1; HSSP; P80362; 1WTL.
                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MU X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBirel. 13, Created)
01-MAY-2000 (TrEMBirel. 13, Last sequence update)
01-DEC-2001 (TrEMBirel. 19, Last annotation update)
Myosin-reactive immunoglobulin light chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
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InterPro; IPRO03596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
NON_TER 1
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3DLINE=98277139; PubMed=9614934;

1411 B., Van der Merwe P.L.,
                                                                                     cetus.",
                                                                                                            "Myosin-reactive autoantibodies in
                                                                                                                                   Young D.C.;
                                                                                                                                                                                                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                               Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      etus. "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99;
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F035028; AAD56264.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
IPR003006; Ig_MHC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
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Primates;
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Primates;
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91.7%;
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Pred. No. 1.9e
2; Mismatches
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                                                                  87:184-192(1998)
                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                            carditis
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Best Local Similarity
Matches 88; Conserv
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Matches 88;
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Best Local
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Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
NON_TER 108 108
                   Q9UL77
Q9UL77;
01-MAY-2000
01-MAY-2000
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NON_TER
SEQUENCE
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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SMART; SM00406; IGv; 1.
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MEDLINE=98277139; PubMed=9614934;
Van der Merwe P.L.,
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01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Myosin-reactive immunoglobulin kappa chain variable
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01-MAY-2000
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HSSP; P80362; 1
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InterPro; IPR003596; Ig_v.
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                                                                                                                      ARFSGSGSGTEFTLTISSLQSEDFAIYHCQQYNSWPPLTFGGGTKVEIK 108
                                                                                                                                DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGS-SFWTFGQGTKVEIK 108
                                                                                                                                                           EIVLTQSPGTLSLSPGERATLSCRASQSVGSSYLAWYQQKPGQAPRLLIYGAFSRATGIP 60
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                                                                                                                                                                                                                                                                                                                                                      Immunol. Immunopathol.
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                                                                                                                                                                                                                                                    109 AA;
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nilarity 80.7%;
Conservative
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108 AA; 11834 MW;
                   (TrEMBLrel.
(TrEMBLrel.
(TrEMBLrel.
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                                                                PRELIMINARY;
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        2MBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
immunoglobulin light chain variable region
                                                                                                                                                                                                                                                                                                                                          AAD56265.1;
                                                                                                                                                                                                                                                     11761 MW;
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Pred. No. 4.2e-40;
8; Mismatches 11;
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Pred. No. 1.3e-39;
                                                                                                                                                                                                                                                                                                                                                      87:184-192(1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                          Mismatches
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Myosin-reactive

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Regult
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              number of hits satisfying chosen parameters:
                                                                   404.5
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Match
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Gapop 10.0 , Gapext 0.5
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                                       70.4
67.4
63.4
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566
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1: sp_archea:*
2: sp_bacteria:*
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                                                                                                                                                                                                                                                                                                                                                                                             sp_archeap:*
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sp_bacteriap:*
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Q920E9
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Q9u178 homo sapien
Q9u183 homo sapien
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Q9u170 homo sapien
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42.6	42.9	43.7	45.8	46.6	46.8	47.7	47.7	48.9	50.2	55.0	55.7	56.6	56.6	57.2	57.8	58.0	58.9	59.3			59.9							
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mm	6 MW; 5P675C52EC7EE197 CRC64; 6%; Score 524; DB 4; Length 109; 4%; Pred. No. 1.6e-49;	Pfam; PF00047; 19; 1. SMART; SM00406; IGv; 1. NON TER 109 109	munol. Immuno 035036; AAD56 0362; IWTL. ; IPR003006;	SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDLINE=98277139; PubMed=9614934; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M., Young D.C.; "Myosin-reactive autoantibodies in rheumatic carditis and normal fetus.";	••	Q901/8; 01-MAY-2000 (TrEMBLrel. 13, Created) 01-MAY-2000 (TrEMBLrel. 13, Last sequence update) 01-DEC-2001 (TrEMBLrel. 19, Last annotation update) Myosin-reactive immunoglobulin light chain variable region	RESULT 1 Q9UL78 PRELIMINARY; PRT; 109 AA.

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NON TER
SEQUENCE
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P01620;
21-JUL-1986
21-JUL-1986
15-JUL-1999
 MEDLINE=02046598; PubMed=6794615;
Andrews D.W., Capra J.D.;
"Amino acid sequence of the varia
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HSSP; P80362; 1WTL
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                       NCBI_TaxID=9606;
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                                                                                      3ukaryota; Metazoa;
4ammalia; Eutheria;
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EXp. Med. 167:840-852(1988).
1- DISEASE: THE PROTEIN IS ONE OF THE SURFACE IMMUNOGLOBULIN M
AUTOANTIBODIES EXPRESSED IN PATIENTS WITH CHRONIC LYMPHOCYTIC
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1-NOV-1990 (Rel.
5-JUL-1999 (Rel.
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in V-III region SIE.
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COMPLEMENTARITY-DETERMINING-3.
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COMPLEMENTARITY-DETERMINING-2.
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Catarrhini;
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SEQUENCE.
MEDLINE-72188439; PubMed-5027703;
MEDLINE-72188439; PubMed-5027703;
Suter L., Barnikol H.U., Watanabe S., Hilschmann N.;
Suter L., Barnikol H.U., Watanabe S., Hilschmann N.;
"Rule of antibody structure. The primary structure of a monoclonal "Rule of antibody structure. The primary structure of a monoclonal immunoglobulin L-chain of kappa-type, subgroup 3 (Bence-Jones protein immunoglobulin L-chain of kappa-type)
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Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

Immunoglobulin V region; Bence-Jones protein.

23 89 BY SIMILARITY.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Hoppe-Seyler's Z. Physiol. Chem. 353:
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- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
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                                                                                                                                       R;Goni, F.R.; Chen,
                                                                                                                                                                                                                     g kappa chain V-III region (Glo)
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                                                      Title: Structural and idiotypic characterization of the Reference number: A30601; MUID:89215279; PMID:2496160; Accession: B30601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Superfamily: immunoglobulin V region; immunoglobulin; Keywords: heterotetramer; immunoglobulin; 16-91/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title: Structural and idiotypic characterization of the L chains of human IgM autoant;Reference number: A30601; MUID:89215279; PMID:2496160;Accession: H30601
                                                                                                                                                                             Species: Homo sapiens (man)
Date: 29-Jun-1989 #sequence
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Property Local Similarity
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                                                                                                                                                            Accession: B30601
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21-129/Product: Ig kappa chain
                                                                                                                   ioni, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Immunol. 142, 3158-3163, 1989
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18-129/Region: J segment (JK1)
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                                                                                                                                                                                                                                                                                                                     DRFSGSGSGTDFTLTTSRLEPEDFAVYYCQQYGSSPYTFGQGTKLEIK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 540; DB 2;
Pred. No. 2.3e-38;
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A; Residues: 1-10
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C; Keywords: hete
F; 16-91/Domain:
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R;Goni, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson, J. Immunol. 142, 3158-3163, 1989

A;Title: Structural and idiotypic characterization of the L chains of human IgM A;Reference number: A30601; MUID:89215279; PMID:2496160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig kappa chain V-III region (Cur) - human (fragment)
C;Species: Homo sapiens (man)
C;Bate: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 21-Jan-2000
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97.2%;
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Pred. No. 3.4e-38;
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Pred. No. 5e-38;
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OM protein - protein search, using sw model

Run on: April 16, 2003, 17:11:36; Search time 33.3474 Seconds (without alignments)
311.344 Million cell updates/sec

Perfect score: Sequence: US-09-644-66BA-7
566
1 BIVLTQSPGTLSLSPGERAT......CQQYGSSPWTFGQGTKVEIK 108

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

Scoring table:

number of hits satisfying chosen parameters:

283224 seqs, 96134422 residues

Minimum DB Maximum DB seq length: 0 length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 73:* 1: pir1:* 2: pir2:*

Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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508.5	509	510	510.5	513	513	515	515	520	524	524	525	525	528.5	529	530	530	531	531	532	532	533	535	536	538	539	540	546	549	Score	,
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ALIGNMENTS

RESULT 1

Ig_kappa chain V-III region (Pie) - human (fragment)

Species:

Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 21-Jan-2000

R;Goni, F.R.; Accession: C30608

Gon1, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson, Immunol. 142, 3158-3163, 1989 D.; S.

A;Accession: C30608 A; Title: Structural and idiotypic characterization of the L chains of human IgM autoa. A; Reference number: A30601; MUID:89215279; PMID:2496160

A;Status: preliminary

A; Molecule type: protein A; Residues: 1-108 < GON>

F;16-91/Domain: immunoglobulin homology Superfamily: immunoglobulin V region; immunoglobulin homology Keywords: heterotetramer; immunoglobulin

<1MM>

Matches Query Match Local 105; Similarity Conservative 97.0%; Score 549; DB 2; Pred. No. 4.1e-39; Mismatches Length 108, Indels 0 Gaps 0,

밁 S 1 EIVLTQSPGTLSLSPGERATLSCRASQSVGSSYLAWYQQKPGQAPRLLIYGAFSRATGIP. 60 ETVLTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATGIP 60

Ś 61 61 DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKVEIK 108 DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGSSPWTFGQGTKLEIK 108

Ig kappa chain precursor V-III region (Hic) - human

Species: Homo sapiens (man)

Accession: PL0021 Date: 30-Jun-1990 #sequence_revision 30-Jun-1990 #text_change 21-Jan-2000

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A; Molecule type: mRNA A; Residues: 1-129 < KIP>

Comment: The protein is

one of the surface immunoglobulin M autoantibodies expressed

A; Cross-references: GDB:136266

A;Gene: GDB:IGKV3 C;Genetics:

A;Map position: 2p12-2p11 C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (k hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into immunoglobulin Cases, sur V region;

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Result
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A; Cross references: EMBI:x68248; NID:g62535; PIDN:CAA48319.1; PID:g62536 A; Note: the nucleotide sequence was submitted to the EMBI Data Library, A A; Note: only a small part of the coding sequence given A; Note: only a small part of the translation is shown R; Njolstad, P.R.; Molven, A.; Apold, J.; Pjose, A. EMBO J. 9; S15-524, 1990 A; Title: The zebrafish homeobox gene hox-2.2; transcription unit, potenti A; Reference number: S08639; MUID:90151628; PMID:1968004 A; A; Accession: S08642
                                                                                                                                                                                                                                                                                                                                                                         R;Molven, A.; Hordvik, I.; Njolstad, P.R.
Biochim. Biophys. Acta 1173, 102-106, 1993
A;Title: Sequence analysis of the zebrafish hox-B5/B6 region.
A;Reference number: S32563; MUID:93250038; PMID:8097929
A;Accession: S32563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N;Alternate names: homeotic protein Hox 2.2
C;Species: Brachydanio rerio (zebra fish)
C;Date: 08-Dec-1993 #sequence_revision 26-May-1995 #text_change 24-Sep-1999
C;Accession: S32563; S08642; S25594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: EMBL:X17267 C;Superfamily: unassigned homeobox proteins; homeobox homology C;Keywords: DNA binding; homeobox; nucleus; transcription regulation
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A; Residues: 1-165 < NJO>
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A;Title: The zebrafish homeobox gene hox-2.2: to
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C;Species: Brachydanio rerio (zebra fish)
C;Date: 21-Nov-193 #sequence_revision 26-May-1995 #text_change 15-Oct-1999
C;Accession: S08641
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A;Residues: 1-228 <MOL>
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9639; MUID:90151628; PMID:1968004
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; Pred. No. 8.1;
1; Mismatches
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A;Status: preliminary; translated fr
A;Molecule type: DNA
A;Residues: 1-72 <GRA>
A;Cross-references: EMBL:U42580; NID
A;Experimental source: specific host
C;Genetics:
A;Gene: a95R
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C,Genetics:
A;Gene: Hox-B6; Hox 2.2
A;Introns: 114/1
C;Keywords: DNA binding; homeobox proteins; homeobox homology
C;Keywords: DNA binding; homeobox; nucleus; transcription regulation
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                                                                                                                                                                                                    Cross-references: EMBL:U42580; NID:94028896; PIDN:AAC96463.1;Experimental source: specific host Chlorella strain NC64A
                                                                                                                                                                                                                                                                                                                                                                              Accession:
                                                                                                                                                                                                                                                                                                                                                                                                               Species: Chlorella virus PBCV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Molecule type: DNA
Residues: 1-603 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana, Reference number: A84420; MUID:20083487; PMID:10617197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.D.; Benito, M.I.; Town, C.D.; Fujii, C.Y.
J. Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, 188, D.; Nierman, M.C.; White, O.; Bisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Les: Arabidopsis thaliana (mouse-ear cress)
Le: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change
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85.7%;
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Library, May
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R;Rieger, M.; Gabel, C.; Mueller-Auer, S.; Schaefer, M.; Zipp, submitted to the Protein Sequence Database, December 1999
                                                hypothetical protein T9C5.40 - Arabidopsis thaliana (5/5pecies: Arabidopsis thaliana (mouse-ear cress) (7/5pate: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change
                                                                                                                                                                                                밁
                                          C;Accession: T46222
                                                                                                                                            RESULT 7
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A;Description: involved in cell morphogenesis interdependently of the C;Keywords: glycosyltransferase; hexosyltransferase; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: T
R;Katayama, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  alpha-glucan synthase (EC 2.4.1.-) mok12 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Gene: arrwoor
C;Superfamily: hypothetical protein b0838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium
A;Reference number: AB1807; MUID:21595285; PMID:11759840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         glutathione S-transfe
C;Species: Nostoc sp.
A;Note: Nostoc sp. st
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                                                                                                                                                                                                                                                                                                                                                                  submitted to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:BA000019; PIDN:BAB76360.1; A;Experimental source: strain PCC 7120
                                                                                                                                                                                                        Status: Procession (Notecule type: DNA ; Molecule type: DNA ; Residues: 1-2352 <KAT>; Cross-references: EMBL:AB018381; PIDN:BAA76558.1 ; Cross-references: EMBL:AB018381; PIDN:BAA76558.1
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Best Local
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                                                                                                                                                                                         Gene: mok12
                                                                                                                                                                                                                                                                                                                  Reference number: Z22509
Accession: T43431
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Species: Schizosaccharomyces
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .; Nakamura, Y.; Wolk, C.P.; Kuritz, T.;
N.; Shimpo, S.; Sugimoto, M.; Takazawa, N
                                                                  Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S-transferase [imported] - Nostoc sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nake, Shimpo, S. 13, 2001
                                                                                                                                                                                                                                                                                                                                                Fission yeast
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ission yeast alpha-glucan
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Pred. No.
                                                                                 Score 30; DB 2;
Pred. No. 1.9e+0;
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glucan synthase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                supersensitive
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                                                                                             Length 2352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 185;
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M.; Yamada, M.; Yasuda, M.; Tabata
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                                                                                                                                            of the actin
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Listing first 45 summaries
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                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        number of hits satisfying chosen parameters:
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Match Length DB
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Copyright (c) 1993 - 2003 Compugen_Ltd.
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KY3B HUMAN
YFUS RAT
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Q9uul4 schizosacch
P02981 escherichia
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homo sapien
homo sapien
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9 mesocricetu
7 mycobacteri
7 prochloroco
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5 human papil
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homo sapien
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4349 4351 108	1088 1333 1374	572 916 933
		<u> </u>
FAT2_HUMAN FAT2_RAT CYC_EISFO YFBJ_SALTI	PIGO_HUMAN YN99_YEAST VCAP_HSV11	BLRO MYRVE MUTS RHIME PRGR HUMAN
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Q9nyq8 O88277 P00030	Q8teq8 P53756 P06491	
homo sapien rattus norv eisenia foe	homo sapien saccharomyc	myrothecium rhizobium m
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DR D	ጟጟጅጟ86,6888888888888	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	RESU HXB6 AC DT DT DT DT OC OC
EMBL; X17267; CAA35171.1; EMBL; X68248; CAA48319.1; PIR; S08642; S08642. PIR; S08642; S08642. PIR; S25594; S25594. PIR; S32563; S32563. HSSP; P02833; HIOM. TRANSFAC; T03664; ZFIN; ZDB-GENE-990415-106; hoxb6a. InterPro; IPR001827; Antennapedia. InterPro; IPR001827; Antennapedia. Pfam; PF00046; homeobox; 1. PRINTS; PR00025; ANTENNAPEDIA. PRINTS; PR00024; HOMEOBOX. ProDom; PD000010; Homeobox; 1.	NE=932500 n A., Hor n A., Hor n A., Hor in A	PubMed=1968004; PubMed=1968004; Hven A., Apold J., Fjose A.; meobox gene hox-2.2: transcription unit, potential s and in situ localization of transcripts."; (1990).	

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Query Match
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A Spouros J., Feat N., Hayles J., Baker S., Basham D., Bowman S.,
Barooks K., Brown D., Brown S., Chillifigworth T., Churcher C.M.,
Collins M., Connor R., Cronin A., Davis P., Feltwell T., Fraser A.,
A Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,
Hallin N., Harris D., Hidalgo J., Hodgson G.,
Hallin N., Harris D., Hidalgo J., Hodgson G.,
Hornoby T., Howarth S., Huckle B.J., Hunt S., Jagels K.,
James K., Jones H., Jones M., Leather S., McDonald S., McLean J.,
Mooney P., Moule S., Mungall K., Murphy L., Niblett D., Odell C.,
A Mooney R., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch B.,
Rutherford K., Outell S., Saunders D., Seeger K., Sharp S.,
Skelton J., Simmonds M., Squares R., Squares S., Stevens K.,
Taylor K., Taylor R.G., Tivey A., Walsh S.V., Warren T., Whitehead S.,
Woodward J., Volckaert G., Aert R., Robben J., Grymonprez B.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Meller-Auer S.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Muller-Auer S.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Muller-Auer S.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Muller-Auer S.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Muller-Auer S.,
Weltjens I., Vanstreels B., Rieger M., Schaefer M., Muller-Auer S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=972;
Katayama S., Dai H., Arellano M., Perez P., Toda T.;
"Fission yeast alpha-glucan synthase Mokl localizes closely with actin and play a role essential for cell morphogenesis and protein kinase C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9UUI4; O13605;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-UNN-2002 (Rel. 41, Last amotation update)
Cell wall alpha-1,3-glucan synthase moki2 (EC MOKI2 OR SPBC3ZH8 13C OR P1011.
                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=20089027; PubMed=10620777;
Machida M., Yamazaki S., Kunihiro S., Tanaka T.,
Haikawa Y., Yamazaki J., Yamamoto S., Sekine M.,
Haikawa Y., Yamazaki J., Kudoh Y., Kikuchi H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA BIND
                                                                                                                                                                                                                                                                                                                        genomic DNA and cDNAs encoded on the segment.";
Yeast 16:71-80(2000).
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Eukaryota; Pungi; Ascomycota; Schlzosaccharomycetes;
                                                                                                                                                                                                                                       MRDLINE=21848401; PubMed≕11859360;
Wood V., Gwilliam R., Rajandream M
                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                        yeast chromosome
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PROSITE; PS00027; HOMEOBOX
PROSITE; PS50071; HOMEOBOX
PROSITE; PS00032; ANTENNAPI
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HOMEOBOX 2; 1.
ANTENNAPEDIA; 1.
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224
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85.7%;
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Pred. No.
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Oguchi A., Nagai
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Matches 6
REVISIONS, AND IDENTIFICATION OF PROTEIN MEDLINE=83263146; PubMed=6307828;
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                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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Bacteria; Proteobacteria;
Escherichia.
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P02981;
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EMBL; AB004534; BAA21388.1;
EMBL; AL590971; CAC37503.1;
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                            REVISIONS,
                                                                                                                          "Directed mutagenesis method
                                                                                                                                                          Livneh
                                                                                                                                                                            MEDLINE=83117828;
                                                                                                                                                                                                      REVISIONS,
                                                                                                                                                                                                                                                                                                       "Complete
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Pfam; PF00534; Glycos_transf_1; 1.
Cell wall; Transferase; Glycosyltransferase.
SEQUENCE 2352.AA; 266561 MW; 78ADF9C2F77
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-i- SIMILARITY: BELONGS TO THE GLYCOSYLTRANSFERASE FAMILY 1.
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-!- CATALYTIC ACTIVITY: UDP-glucose +
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1-JUL-1986 (Rel.
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                                                                      utagenesis method for analysis of mutagen specificity: to ultraviolet-induced mutagenesis.";
Acad. Sci. U.S.A. 80:237-241(1983).
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resistance protein, class C (TETA(C)).
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28; PubMed=6337373;
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Copyright (c) 1993 - 2003 Compugen Ltd.
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O91uu6 escherichia
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O22787 arabidopsis
Q84416 paramecium
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                                                                                                                                                                                                                                  Ribeiro S.G., Ambrozevicius L.P., de Avila A.C., Calegario R.F., Fernandes J.J., Lima M.F., Mello R.N., Rocha H., Zerbini F.M.; "Distribution and genetic diversity of tomato-infecting geminiviruses
                                                                                                                                                                                                                                                                                                      Tomato infectious yellows virus.
Viruses; ssDNA viruses; Geminiviridae; Begomovirus
NCBI_TaxID=172393;
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                    SEQUENCE FROM N.A.
MEDLINE=20478054; PubMed=11021991;
Sun L., Gurnon J.R., Adams B.J., Graves M.V.,
"Characterization of a beta-1,3-glucanase enc
                                                                                                                                                                                    SEQUENCE FROM N.A.

MEDILINE=95407089; PubMed=7676624;

Li Y., Lu Z., Burbank D.E., Kutish G.F., F
"Analysis of 43 kb of the Chlorella virus
positions 45 to 88.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-20083467; PubMed=10617197;
Lin X., Kaul S., Rounsley S.D., Shea T.P., Benito M.-I., Town C.D., Pujii C.Y., Mason T.M., Bowman C.L., Barnstead M.E., Feldblyum T.V., Buell C.R., Ketchum K.A., Lee J.J., Ronning C.M., Koo H., Moffat K.S., Cronin L.A., Shen M., Venaken S.E., Umayam L., Tallon L.J., Gill J.E., Adams M.D., Carrera A.J., Creasy T.H., Goodman H.M., Somerville C.R., Copenhaver G.P., Preuss D., Nierman W.C., White O., Eisen J.A., Salzberg S.L., Fraser C.M., Venter J.C.;
"Sequence and analysis of chromosome 2 of the plant Arabidopsis
                                                                                                                                                                                   Positions
Virol
                                                                                                                                                                                                                                                         Paramecium bursaria chlorella virus 1
!Iruses; dsDNA viruses, no RNA stage;
!NCBI_TaxID=10506;
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01-DEC-2001
                                                                                                      MEDLINE=20013326; PubMed=10544099;
Kaiser A., Vollmert M., Tholl D., Graves M.V.,
Lisec A.D., Nickerson K.W., Van Etten J.L.;
"Chlorella virus PBCV-1 encodes a functional ho
                                                                                                                                                                                                                                                                                                A95R.
                                                                                                                                                                                                                                                                                                         A95R protein.
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STRAIN=CV. COLUMBIA;
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Bukaryota; Viridiplantae; Streptophyta; Embry.
Spermatophyta; Magnoliophyta; eudicotyledons;
eurosids II; Brassicales; Brassicaceae; Arabi
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AT2G33360.
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     276:27-36 (2000)
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DREAD REPRESENTATION OF THE PROPERTY OF THE PR
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ID Q9
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                                                                A Kawai J. Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,

A Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,

A Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,

RA, Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,

A Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,

RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,

RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,

RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,

RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,

RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,

RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,

RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,

RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,

RA Nordone P., Raing B., Ringwald M., Rodriguez I., Sakamoto N.,

RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,

RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,

Havashitaki F., Sokamoto N., Kawaji H., Kohtsuki S.,
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2210404M20RIK.
Mus musculus (Mouse).
Mus musculus (Mouse).
Chordata; C:
Chordata; C:
Chordata; Rodentia; S:
                Nature
                  "Functional annotation of a Nature 409:685-690(2001).
                                                              Hayashizaki Y.;
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Submitted
[5]
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01-JUN-2001
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SEQUENCE FROM N.A.
Graves M.V., Van Etten J.L.;
Graves M.V., Par Etten J.L.;
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Submitted (DEC-2000) to the
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Adachi J., Func...
An S., Yamanaka
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                                  mouse cDNA collection.";
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iya M., Lee N.H.,
Mombaerts P.,
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Gapop 10.0 , Gapext 0.5
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34
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l: /SIDS2/gcgdata/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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/SIDS2/gcgdata/geneseq/
                                                                                                                                                                                                                                                 SIDS2
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4	44	44	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
27	27	27	27	28	28	28	28	28	28	28	28	28	28	28	28	28	28	28	28	28	28	28	29	29	29	29	29	29	29	29	29	29	29	29
79.4	79.4	79.4	79.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	82.4	•	82.4	82.4	85.3	85.3	85.3	85.3	85.3	85.3	85.3	85.3	85.3	85.3	85.3	85.3
95		28		2139	924	832	820	808	791	724	675	618	587	587	524	410	361	287	259	179	143	85	1298	415	396	396	396	. 396	396	373	359	177	177	61
23	21	17	22	22	22	22	21	22	22	23	22	22	23	22	21	20	23	21	22	22	22	22	21	22	23	22	20	20	19	22	22	21	21	22
ABP02653	AAB40534	AAR91383	AAM97727	ABB62393	AAU09958	AAB74457	AAB43354	AAB65679	AAE11780	ABB04802	AAM40778	AAB74769	ABG65328	AAB74754	AAB58954	AAW93254	ABB49841	AAB35708	ABG27325	AAU53900	AAU56695	AAU53827	AAB53204	ABB52991	AAU75552	AAB50899	AAY15383	AAY42545	AAW56695	ABG01757	AAB30608	AAB51686	AAB51685	AAU41549
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ORF	-	ME#1 library deriv	Human peptide #100	phila	cadher			protein	Human kinase (PKIN	LDL receptor bindi	Human polypeptide	Human secreted pro				Human ESRP1 protei	Listeria monocytog	Human protein cont	Novel human diagno	Propionibacterium	Propionibacterium	Propionibacterium	3	richia coli	celZ integration v	encoded	Plasmid pRZTL1, Te	٠.	Tetracycline resis		Amino acid sequenc	Human secreted pro	Gene 22 human secr	Propionibacterium

AAB67494;

29-MAY-2001

(first entry)

AAB67494 standard; peptide; 7 AA

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RESULT 1
AAB67494
ID AAB67494
XX AAB674
AC AAB674
XX DT 29-MAY
XX COmpl(
KW Coylot,
KW Cownlti
KW rheum
XX DO 01-M
XX DO -M
XX MEI
XX MEI
XX MEI
XX WPI
                                                                                                                                                                                                                                                                      autoimmune disease; infectious disease; rheumatoid arthritis; myasthenia gravis; multiple sclerosis; insulin-dependent distransplant rejection; graft versus host distransplant rejection;
                                               Korman AJ, Halk EL,
                                                                                                                 24-AUG-1999;
                                                                                                                                             24-AUG-2000; 2000WO-US23356.
                                                                                                                                                                            01-MAR-2001.
                                                                                                                                                                                                           WO200114424-A2
                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                   Complementarity determining region; CDR; immune response; antibody; cytotoxic T lymphocyte associated antigen-4; CTLA-4; B7 ligand; can
                                                                                                                                                                                                                                                                                                                                                                                 Human light chain complementarity determining region 2 (CDR2).
                                                                                (MEDA-) MEDAREX INC.
                                                                                                               99US-0150452.
                                                 Lonberg
                                                  z
                                                                                                                                                                                                                                                                                     diabetes mellitus; inflammation,
                                                                                                                                                                                                                                                                                                                    inflammation; allergy;
                                                                                                                                                                                                                                                                                                                                ligand; cancer;
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Result

Query Match Length DB

ä

Description

Human light chain Anti-HIV gp41 imm

SUMMARIES

100.0

AAB67494 AAR54279

AAY98283 AAY34471

Anti-HIV gp41 immu
VI of Fab, SS 41 8
Anti-gp41 light ch
Anti-gp41 light ch
Porphorymonas ging
Porphorymonas ging
Porphorymonal ging
Novel human diagno
Novel human diagno
Novel human diagno

WPI; 2001-202933/20

AAY95174 AAW0132

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RESULT 2
AAR54279
ID AAR5
XX AAR5
AC AAR5
XX AAR5
AC AAR5
XX AIII
XX Huma
XX Huma
XX Homo
XX H
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ~Yuman immunodeficiency virus; HIV1; glycoprotein gp41; epitope;
jutralisation; monoclonal antibody; light chain; variable region;
juramework region; complementarity determining region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibody preparation comprising two antibodies of the invention are useful for suppressing a immune response in a patient. They are used for treating cancer, infectious diseases and promoting beneficial autoimmune creations for the treatment of diseases with inflammatory or allergic omponents. The polyvalent or polyclonal preparations are useful for creating autoimmune diseases such as rheumatoid arthritis, myasthenia gravis and lupus erythematosus, multiple sclerosis, insulin-dependent diabetes mellitus, transplant rejection, and inflammation, graft versus
                                                                      W09497922-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-HIV gp41 immunoglobulin light chain V region clone SS 41 8.
                               14-APR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     also useful for treating autoimmune disease in a subject caused or exacerbated by increased activity of T cells and for treating prostate cancer, melanoma or epithelial cancer. A polyvalent or polyclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibodies are used in methods for inducing, augmenting or prolonging an immune response to an antigen in a patient, where the antibodies block binding of human CTLA-4 to human B7 ligands. The antibodies are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB67490-99 and AAB67501-06 represents complementarity determining regions (CDRs) of human antibodies. The antibodies specifically bind to human cytotoxic T lymphocyte associated antigen-4 (CTLA-4). Such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human sequence antibody that binds to human cytotoxic T
lymphocyte associated antigen-4, useful for inducing, augmenting
prolonging immune response to antigen or for suppressing immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human cytotoxic T ly antibodies are used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 30; Page 99; 127pp; English.
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168 7; Conserv
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ilarity 100.0%;
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                                                                                                                                                                                          label= FR3
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                                                                                                                                                                                                                                                                           abel= FR2
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Pred. No. 7.8e+05;
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Matches
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Best Local (
WO9602273-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lymphocyte mRNA was converted to cDNA and subjected to PCR amplification using primers specific for heavy and light chain variable regions. The amplification products were inserted into a dicistronic vector to produce a library of fragments. E.coli Xii Blue cells were transformed with the library. Filamentous phage were produced which expressed the MAB regions on their surface. Panning with gpl20 and gp41 resulted in the recovery of immunoreactive clones. The light chain VL region sequence AARS4279 neutralises HIV1
                                                 Region
                                                                                                                     Region
                                                                                                                                                      Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW01324 standard; Protein; 112
                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; Page 218; 248pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-135516/16.
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1; Mismatches
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A
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Listing first 45 summaries
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Perfect score:
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Match Length
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Gapop 10.0 , Gapext
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l: /cgn2_6/ptodata/1/pubpaa
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US-09-838-529-2
US-09-838-708-45
US-09-828-708-45
US-09-828-708-48
US-09-828-708-48
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US-09-828-708-48
US-09-828-708-84
US-09-828-708-84
US-09-828-708-86
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                           Sequence 138, App
Sequence 62, App
Sequence 2, Appl
Sequence 240, App
Sequence 107, App
Sequence 47, Appl
Sequence 48, Appl
Sequence 48, Appl
Sequence 2362, Appl
Sequence 2362, Appl
Sequence 23, Appl
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٠	US-10-180-557-562	•	US-10-176-913-562	57-	'n	US-10-175-752-562	6	06-56	63-	75-737-56	US-10-063-616-166	58-56	US-10-174-590-562	US-10-063-547-166	US-09-880-748-1499	US-09-815-242-12053	US-09-880-748-1219	US-09-880-748-1035	US-09-880-748-1264	US-09-880-748-1896	ö	US-09-859-053-34	US-09-815-242-10606	US-10-001-934-40	US-10-025-687-10	US-10-125-687-10
	562,	æ	e 562,	æ	e 562,	e 562,	Sequence 562, App	e 562,	e 166,	æ	e 166.	e 562,	æ	166,	-	ce 12053,	e 1219, A	Sequence 1035, Ap		1896,	ы	34.	e 106	40.	Sequence 10, Appl	Sequence 10, Appl

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RESULT 2
US-10-102-806-662
; Sequence 662, Application US/10102806
; Publication No. US20030054421A1
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                                                                                                                                                                                                                                            ; OTHER INFORMATION: fungal gene US-10-029-180-138
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                                                                                                                                                                                      Query Match
Best Local S
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CURRENT FILING DATE: 2001-12-22
PRIOR APPLICATION NUMBER: US 60/257,431
PRIOR FILING DATE: 2000-12-22
NUMBER OF SEQ ID NOS: 138
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 138, A Publication No.
                                                                                                                                                                            Matches
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TITLE OF INVENTION: No. US20020182708A1el
FILE REFERENCE: MIC-004
                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                           LENGTH: 805
                                                                                                           797 GAWSRAT 803
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6; Conserv
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Milna, G. Todd
Sherman, Amir
Silva, Jeffry C.
Trueheart, Josh
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Madden, Kevin
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                                                                                                                                                                          Conservative
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                                                                                                                                                                                      85.3%;
85.7%;
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Pred. No. 1.4e+02;
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Thes 5; Conserv:
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US-09-838-529-2
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                                            Sequence 240, Application US/09893737 Patent No. US20020110855A1
                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 924
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/838,529
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/US00/05881-
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: Curtis, Rory A. J.

ITLE OF INVENTION: 57809 AND 57798, NOVEL HUMAN CADHERIN
ITLE OF INVENTION: MOLECULES AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: SITE
LOCATION: (191)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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THER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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E OF INVENTION: Nucleic Acids, Proteins and Antibodies
REFERENCE: PA103PIC1
                                                                                                                                 GSYSRAT 39
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                                 INFORMATION
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                               ; • ORGANISM: Pseudomonas aeruginosa
US-09-975-719-107
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US-09-893-737-240
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                                   CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2001-10-10
PRIOR APPLICATION NUMBER: US 09/199,637
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: US 60/066,517
PRIOR FILING DATE: 1997-11-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 437
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PRIOR FILING DATE: 2000-06-30
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                                                                                                                    PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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                                                               TLE OF INVENTION: Autoantibodies to glucosé-6-phosphate isomerase TLE OF INVENTION: autoimmune disease LE REFERENCE: 1361.005US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
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                                                                                                                                                                                                                                                                                         127 GAFEKAT 133
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LE OF INVENTION: SEQUENCES AND USES THEREOF
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Schaller,
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5. US20030022349A1
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Pred. No. 2e+02;
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Gapop 10.0 , Gapext 0.5
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     Issued Patents AA:*
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/cgn2_6/ptcdata/1/1aa/5B_COMB.pep:*
/cgn2_6/ptcdata/1/1aa/6A_COMB.pep:*
/cgn2_6/ptcdata/1/1aa/6A_COMB.pep:*
/cgn2_6/ptcdata/1/1aa/6CTUS_COMB.pep:*
/cgn2_6/ptcdata/1/1aa/backfiles1.pep:*
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US-08-850-880-4
US-08-814-877-4
US-09-272-432A-4
US-09-272-432A-4
US-09-199-637A-107
US-08-162-102C-39
US-08-112-102C-39
US-08-112-788A-14
US-08-477-531B-14
US-08-477-531B-14
US-08-477-531B-14
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US-08-488-113B-150
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US-08-232-081B-42
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CORRESPONDENCE ADDRESS:	
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: TITLE OF INVENTION: TO HIMAN IMMINISTRATICIENCY VIRIS	
. ADDITONT: Lerner Dichard A	
: burcon, Demins	
MMATION:	
C.F	
TTD_0D_076_0E0_1E1	

Conservative

Score 31; DB Pred. No. 4.6; 1; Mismatches

DB 1;

Length 112;

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Gaps

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               US-08-899-575-151
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Best Local Similarity
""" hes 6; Conserve
Sequence 151, Application US/08899575
                                                                                                                                   Best Local Similarity
Matches 6; Conserv
                                                                                                                                                              Query Match
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                       OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: SC
TELECOMMUNICATION INFORMATION:
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                                                                          51 GAYSRAT 57
                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER:
                                                                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
                                                                                                       1 GAFSRAT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CATION NUMBER: US/08/899,575
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                                                                                                                                                                                                                                                                                                                                                                                  AGENT INFORMATION:
                                                                                                                                                                                                                                                112 amino acids
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                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jolla
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                             619-554-2937
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566 No. 5770440th Torrey Pines Road, Suite 220,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The Scripps Research Institute, Office of
                                                                                                                                                                                                                                                                                                                                                                                                                                            30-SEP-1993
                                                                                                                                                                                                       protein
                                                                                                                                                                                                                                                                                                                                                                                                 JMBER: US 07/954,148
30-SEP-1992
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18-JUL-1994
JMBER: US 08/178,302
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                                                                                                                                                                                                                                                                                                                                                     34,163
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                                                                                                                                             Score 31;
Pred. No.
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Query Match
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"arches 6; Conserv:
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                                                                                                                                                                   equence 151, Application PC/TUS9508743 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ELECOMMUNICATION INFORMATION:
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FURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                       51 GAYSRAT 57
                                                                                                                   TLE OF INVENTION:
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APPLICATION NUMBER: US 07/954,148
FILING DATE: 30-SEP-1992
                                                                                 MBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                          1 GAFSRAT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/899,575
                             EDIUM TYPE: Floppy disk
OMPUTER: IBM PC compatible
PERATING SYSTEM: PC-DOS/MS-DOS
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£: CA
TRY: USA
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RE: PatentIn Release #1.0, Version #1.25 (EPO)
APPLICATION DATA:
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JMBER: US 08/178,302
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170
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                                                                                                                                 HUMAN NEUTRALIZING MONOCLONAL ANTIBODIES
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Pred. No.
                                                                                                                   IMMUNODEFICIENCY VIRUS
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed.
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derived by analysis of the total
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Match Length
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Gapop 10.0 ,
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93
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71	72	74	74	75	75	75	75	75	75	75	77	77	78	78	82
76.3	77.4	79.6	79.6	80.6	80.6	80.6	80.6	80.6	80.6	80.6	82.8	82.8	83.9	83.9	88.2
119	121	118	110	133	128	122	119	118	114	108	137	122	133	76	139
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RESULT 1

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Ig heavy chain V region (clone alpha-TNF-A1) - human (fragment)
C;Species: Homo mapiens (man)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 23-Jul-1999
C;Accession: S36259
R;Griffiths, A.D.; Malmqvist, M.; Marks, J.D.; Bye, J.M.; Embleton, M.J.; McCafferty, JEMBO J. 12, 725-734, 1993
A;Title: Human anti-self antibodies with high specificity from phage display libraries.
A;Reference number: S36256; MUID:93178448; PMID:7679990
A;Accession: S36259
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S36259
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                                                                                                                                                                          A;Cross-references: EMBL:Z18850; NID:g33123; PIDN:CAA79302.1; PID:g939902 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-98 <TOM>
A;Residues: 1-98 <TOM>
A;Crose-references: EMBL:Z17389; NID:g32835; PIDN:CAA78994.1; PID:g32836
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig heavy chain V region (COS 3) - human (fragment)
c;Species: Homo sapiens (man)
C;Date: 07-Jan-1994 #sequence_revision 17-Nov-1995 #text_change 23-Jul-1999
c;Accession: S29543
R;Tomilinson, M.; Walter, G.; Cook, G.P.; Winter, G.
submitted to the EMBL Data Library, October 1992
A;Reference number: S29543
A;Accession: S29543
A;Accession: S29543
                                                                                                                                                                                                                                                                         A;Status: preliminary; nucleic acid sequence not shown A;Molecule type: mRNA A;Residues: 1-117 <GRI>
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                       1 FISYDGNNKYYADSVKG
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FIRYDGSNKYYADSVKG 66
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Similarity 88.2%;
15; Conservation
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                                                                                     Score 83; DB 2;
Pred. No. 1.1e-05;
1; Mismatches 1
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Pred. No. 9.2e-06;
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A; Reference number: $70442; M
A; Accession: $70442
A; Status: ot compared with c
A; Molecule cype: mRNA
A; Residues: $-140 <CUI>
C; Superfamily: immunoglobulin
                                                                                                                                          C;Species: Homo sapiens (man)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change C;Accession: S70442
C;Accession: A.M.; Fumoux, F.; Fougereau, M.; Tonnelle, C.
Mol.; Immunol. 29, 1363-1373, 1992
                                                                                        Mol. Immunol. 29, 1363-1373, 1992
A;Title: IgM kappa/lambda EBV human B cell clone: an early step of differentiation
A.Pefarence number: S70442; MUID:93024508; PMID:1383695
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Acate: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31688
C;Accession: S31688
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
                                                                                                                                                                                                                                                         Ig_heavy chain precursor V region (mu) - human (fragment)
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R; Cuisinier, A.M.;
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A;Residues: 1-134 <CUI>
A;Residues: 1-134 <CUI>
A;Cross-references: EMBL:Z14174; NID:g31013; PIDN:CAA78543.1; PID:g31014
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
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submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Reference number: S31585
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ∞¥y Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    preliminary
                                                                 not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                               FIRYDGSNKYYADSVKG
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15; Conserv
  immunoglobulin V region;
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Pred. No. 1.3e-05;
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immunoglobulin homology
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1.3e-05;
1;
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                                                            Ig heavy chain V-III region (AW-Vx) - human (fragment)
                                                                                                     RESULT
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A; Accession: S44115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Experimental source: B cells from patient TD with acute lymphoblastic l A;Note: the sequence shows the V region (TD-Vo) from a nonproductive DNA C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin
                                                                                     PL0116
                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-97 < HAW>
                                                                                                                                                                                                                                                                                                                                                                                                                             A; Description: Idiotypic
                                                                                                                                                                                                                                                                                                                                                                                                                                           submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: mRNA
A; Residues: 1-94 <BIR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Bird, J.; Galili
J. Exp. Med. 168,
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J heavy cma...
;Species: Homo sapiens (man,
;Date: 07-Jun-1990 #sequence_revision
;Date: 07-Jun-1990 #sequence_revision
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C.Species: Homo sapiens (man)
C.Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 16-Aug-1996
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Best Local :
                                                                                                                                                                                                                                                                                               Cross-references: EMBL:231384; NID:9472969; PIDN:CAA83259.1; PID:9940526 Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                          Status: preliminary
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93.8%;
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93.8%;
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Pred. No. 1.3e-05;
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Pred. No. 1.3e-05;
1; Mismatches 0
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Pred. No. 1.3e-05
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                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                        21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 heavy chain V-III region GA.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
SEQUENCE.
MEDLINE=74175307; PubMed=4208843;
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HSSB; P01772; 2PB4.
InterPro; IPR003006; Ig_W:
InterPro; IPR003596; Ig_V:
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
Immunoglobulin V region.
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"BLINES TOND.
"BLINES TOND.
"Rule of antibody structure. The primary structure of a monoclon ratio of antibody structure. The primary structure of a monoclon log1 immunoglobulin (myeloma protein Nie), I: Purification and characterization of the protein, the L- and H-chains, the cyanogen bromide cleavage products, and the disulfide bridges.", Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540 (1976).
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Ponsting1 H., Hilschmann N.;
Ponsting1 H., Hilschmann N.;
"The rule of antibody structure. The primary structure of monoclonal IgG1 immunoglobulin (myeloma protein Nie). I chymotryptic peptides of the H-chain, alignment of the peptides and discussion of the complete structure.";
Hoppe-Seyler's Z. Physiol. Chem. 357:1571-1604(1976).
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 heavy chain V-III region NIE.
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Mammalia; Eutheria;
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1g heavy chain V region 5A precursor.
Carassius auratus (Goldfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cyprinidae; Carassius.
MCBI_TaxID=7957;
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      Proc. Natl. Acad. Sci. U.S.A.
PIR; B28966; B28966.
HSSP; P01772; 2FB4.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
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MEDLINE=8814476; PubMed=3125551;
Wilson M.R., Middleton D., Warr G
"Immunoglobulin heavy chain varia
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Putnam F.W., Liu Y.-S.V., Low T.L.K.;
"Primary structure of a human IgA1 immunoglobulin. IV. Streptococcal
IgA1 protease, digestion, Fab and Fc fragments, and the complete
amino acid sequence of the alpha 1 heavy chain.";
This is a sequence of the alpha 1 heavy chain.";
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable region
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HSSP; P01772; 2FB4.
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Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
Young D.C.;
Myosin-reactive autoantibodies in rheumatic carditis and normal
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                                                                                                                                                                   Score 83; DB 4;
Pred. No. 1.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                            ED57FDD19086D07F CRC64;
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                                                                                                                                      i; Indels
                                                                                                                                                                                                    Length 113;
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09UL93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 15
        Query Match
Best Local /
                                                                                                                         InterPro; IPR003596; IQ
Ffam; PP00047; Ig; 5.
SMART; SM00409; IG; 2.
SMART; SM00407; IGC1; 4.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QBWUK1 ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9UL93;
Q9UL93;
01-MAY-2000
                                                              PROSITE; PS00290; IG_MHC; UNKNOWN_3.
Hypothetical protein.
SEQUENCE 613 AA; 67296 MW; 60C71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                      Strausberg R.;
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=98277139; PubMed=9614934; Wu X.. Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                 nterPro,
                                                                                                                                                                                                                                                     nterPro;
                                                                                                                                                                                                                                                                                         MBL; BC020240; AAH20240.1;
                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                   CBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Young D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Myosin-reactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                        ISSUE=TONSIL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    omo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-MAR-2002 (TrEMBLrel. 20, Cr
1-MAR-2002 (TrEMBLrel. 20, La
1-JUN-2002 (TrEMBLrel. 21, La
YPothetical 67.3 kDa protein.
    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 ISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rPro; IPR003006; Ig_w.
rPro; IPR003596; Ig_v.
; PF00047; ig; 1.
rr; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISYDGSNKYYADSVKG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P01772; 2FB4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunol.
  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                     ; IPR003599; Ig.
; IPR003597; Ig_c1.
; IPR003006; Ig_MHC.
; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  116 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TrEMBLrel.
(TrEMBLrel.
(TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunopathol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBLrel. 13, Created)
EMBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD56257.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     116
12434 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata;
Primates;
                                                            67296 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata;
  88.2%;
93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.2%;
93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence up
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
Score 82; DB 4;
Pred. No. 0.00017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87:184-192(1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                          60C7F5950671E315 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Craniata; Vertebrata; I
Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0DA0348154DD6061 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         613 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    $
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    update)
                 Length 613;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S.X
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RESULT 5
Q9Y509
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                                                                                     myeloma using PCR with patient-specific immunoglobulin gene primers.", Leukemia 9:1948-1953(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9UL84;
01-MAY-2000
01-MAY-2000
                                                                                                                                                                                                                                                                                Homo sapiens (Human).
Bukaryota; Metazoa; Chordata;
Mammalia; Butheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1999
01-DEC-2001
                                                         EMBL; S80860; AAD14339.1;
HSSP; P01772; 2FB4.
                                                                                                                                                           Cao J., Vescio R.A., Rettig M.B.,
Lichtenstein A.K., Berenson J.R.,
                                                                                                                                                                                               MEDLINE=96071149; PubMed=7475288;
                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          605360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AF035030;
HSSP; P01772; 2
                 InterPro; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                   VH3 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Myosin-reactive
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MEDLINE=98277139; PubMed=9614934;

Tin B Van der Merwe P.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalla; Butheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR003006; Ig_WHC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oung D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fragment)
                                                                                                                                                                                                                     QUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 ISYDGNNKYYADSVKG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       122
122 AA;
                                                                                                                                                                                                                                                                                                                                                                                    (TrEMBLrel. 12, (TrEMBLrel. 12, (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        74.2%;
llarity 81.2%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD56266.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13579 MW;
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                                                                                                                                                                                                                                                                                                                                                                                Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 69; DB 4;
Pred. No. 0.003;
2; Mismatches
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Last annotation update)
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                                                                                                                                                                                                                                                                                Craniata; Vertebrata;
Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36054D41366545B8 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                            Hong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatic carditis and normal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     147
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                                                                                                                                                                              C.H.,
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                                                                                                                                                                                                                                                                                  Hominidae;
                                                                                                                                                                              Ki a
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                                                                                                                                                                              A., Lee
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                                                                                                                                                                                                                                                                                                   Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Euteleostomi; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            region
                                                                                                                                                                              J.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
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Perfect score:
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Maximum DB
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                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 number of hits satisfying chosen parameters:
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seq length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       April 16, 2003, 16:56:17 ; Search time 13.4807 Seconds (without alignments) 168.037 Million cell updates/sec
                                                                                                                                                                                                             Match
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93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     908470 seqs, 133250620 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 FISYDGNNKYYADSVKG 17
                                                                                                                                                                         100.0
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/SIDS2/gcgdata/geneseq/geneseqp-embl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GenCore version 5.1.4 p5 4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                         /SIDS2/gcgdata/geneseq/geneseqp-embl/AA2000.
/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2001.
/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2002
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                                                       AAB39830
AAU08361
AAU08384
AAB39836
                                                                                                                                                  AAB67491
AAB67508
                                                                                                                                    AB67513
                                                                                                                                                                                                                                                                    SUMMARIES
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                                                                                                          Human heavy chain
Light chain variab
Heavy chain variab
Anti-hIL12 antibod
                                 Antibody heavy cha
Anti-OPGbp antibod
Anti-hIL12 antibod
Anti-hIL12 antibod
                                                                                                                                                                                                         Description
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antibod
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1-IL12	4	21	115	9	83	5
i-IL12	AAB39547	21	115	9	83	44
1-IL12	954	21	115	٩.	83	43
1-IL12	954	21	115	۰.	83	42
i-IL12	954	21	115	9.	83	41
1-IL12	953	21	115	٠.	83	40
1-IL12	953	21	115	۰.	83	39
1-IL12	AAB39535	21	115	٩.	83	38
1-IL12	953	21	115	٩.	83	37
1-IL12	AAB39531	21	115	89.2	83	36
1-IL12	AAB39529	21	115	٠	83	35
1-IL12	AAB39527	21	115	9	83	34
۳	AAB39525	21	115	9	83	33
1-IL12	952	21	115	9.	83	32
1-IL12	AAB39521 ·	21	115	9.	83	31
1-IL12	AAB39519	21	115	٠	83	30
1-IL12	951	21	115	٠	83	29
1-hIL-1	AAB39515	21	115	9	83	28
i-hIL-12	AAB39507	21	115	9	83	27
1-hIL-12	AAB39499	21	114	٠	83	26
i-hIL12	AAB40141	21	98	۰.	83	25
	AAB40123	21	96	9	83	24
an heavy ch	AAB67496	22	17	٠	83	23
i-hIL12	AAB39881	21	17	٥	83	22
i-hIL12 ant	· AAB39871	21	17	٥	83	21
i-hIL12 ant	AAB39865	21	17	9.	83	20
i-hIL12	AAB39842	21	17	٩	83	9
i-hIL12	AAB39834	21	17	٩	83	18
i-hIL12	AAB39832	21	17		83	17
1-hIL12 a	98	21	17	89.2	83	16
1-hIL-12	951	21	17	٩	83	15
i-hIL-12	950	21	17	۰.	83	14
i-hIL-12	948	21	17	٩	83	13
1-hIL12	ω	21	17		84	12
Anti-hIL12 antibod	w	21	17	ė.	84	11

RESULT 1 AAB67491 Complementarity determining region; CDR; immune response; antibody; cytotoxic T lymphocyte associated antigen-4; CTLA-4; B7 ligand; cancer; autoimmune disease; infectious disease; inflammation; allergy; rheumatoid arthritis; mysathenia gravis; lupus erythematosus; multiple sclerosis; insulin-dependent diabetes mellitus; inflammation; Korman AJ, Halk EL, 24-AUG-1999; 24-AUG-2000; 2000WO-US23356. 01-MAR-2001. WO200114424-A2 Human heavy chain complementarity determining region 2 (CDR2). AAB67491; AAB67491 standard; peptide; 17 AA. Homo sapiens. transplant rejection; 29-MAY-2001 (MEDA-) MEDAREX INC (first entry) 99US-0150452 Lonberg N;

WPI; 2001-202933/20

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RESULT 2
AAB67508
ID AAB6
PXRXBXBXBXBXBXFFFFFFXRXBXBXBXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC AAB67490-99 and AAB67501-06 represents complementarity determining CC regions (CDRs) of human antibodies. The antibodies specifically bind to CC human cytocoxic T lymphocyte associated antigen-4 (CTLA-4). Such CC antibodies are used in methods for inducing, augmenting or prolonging CC an immune response to an antigen in a patient, where the antibodies are CC block binding of human CTLA-4 to human B7 ligands. The antibodies are CC calso useful for treating autoimmune disease in a subject caused or CC exacerbated by increased activity of T cells and for treating prostate CC cancer, melanoma or epithelial cancer. A polyvalent or polyclonal CC antibody preparation comprising two antibodies of the invention are CC treating cancer, infectious diseases and promoting beneficial autoimmune expenses and promoting beneficial autoimmune comprising a manune response in a patient. They are used for CC treating cancer, infectious diseases with inflammatory or allergic components. The polyvalent or polyclonal preparations are useful for comparation autoimmune diseases with a patient or useful for comparation and lupus erythematosus, multiple sclerosis, insulin-dependent compared the propagation, graft versus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                              Region
                          24-AUG-1999;
                                                          24-AUG-2000,
                                                                                            01-MAR-2001.
                                                                                                                          WO200114424-A2
                                                                                                                                                                                                                                                                                                                                            Attoroxic T lymphocyte associated antigen-4; CTLA-4; B7 ligand autoimmune disease; infectious disease; inflammation, allergy; rheumatoid arthritis; myasthenia gravis; lupus erythematosus; multiple sclerosis; insulin-dependent diabetes mellitus; inflammations
                                                                                                                                                                                                                                                                                                                                                                                                                                                Light chain variable region of anti-CTLA-4 antibody 10D1.
                                                                                                                                                                              Region
                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                   Key
                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                    transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB67508 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                Complementarity determining region; CDR; immune response; Ytotoxic T lymphocyte associated antigen-4; CTLA-4; B7 li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human sequence antibody that binds to human cytotoxic T lymphocyte associated antigen-4, useful for inducing, augmenting prolonging immune response to antigen or for suppressing immune response in patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 30; Page 99; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 FISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 FISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                rejection; graft versus host disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
ilarity 100.0%;
Conservative 0;
                                                        2000WO-US23356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                        9908-0150452
                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                          /note=
                                                                                                                                                                                                                            note= "CDR1"
                                                                                                                                                              note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide;
                                                                                                                                                                              107
                                                                                                                                                              "CDR3!
                                                                                                                                                                                            "CDR2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 93;
Pred. No.
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augmenting
                                                                                                                                                                                                                                                                                                                                                                                              ligand; cancer;
                                                                                                                                                                                                                                                                                                                                            inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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                                                                                                                                                            The present sequence represents the light chain variable region of CC human antibody 1001. This antibody specifically binds to human CC cytotoxic T lymphocyte associated antigen-4 (CTLA-4). Such antibodies CC are used in methods for inducing, augmenting or prolonging an immune CC response to an antigen in a patient, where the antibodies block CC binding of human CTLA-4 to human B ligands. The antibodies are CC also useful for treating autoimmune B ligands. The antibodies are CC exacerbated by increased activity of T cells and for treating prostate CC cancer, melanoma or epithelial cancer. A polyvalent or polyclonal CC antibody preparation comprising two antibodies of the invention are CC useful for suppressing a immune response in a patient. They are used for treating cancer, infectious diseases and promoting beneficial autoimmune CC reactions for the treatment of diseases with inflammatory or allergic components. The polyvalent or polyclonal preparations are useful for treating autoimmune diseases such as rheumatoid arthritis, myasthenia CC gravis and lupus erythematosus, multiple sclerosis, insulin-dependent const disease.
                                         Matches
                                                                            Query Match
                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 25; Fig
                                                                                                                                                          host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human sequence antibody that binds to human cytotoxic lymphocyte associated antigen-4, useful for inducing, augmen prolonging immune response to antigen or for suppressing imm response in patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Korman AJ,
                                 17; Conserv
                                                                                                                   118 AA;
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                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7; 127pp;
                                                     100.0%;
                                     0
                                                     Score 93;
Pred. No.
                                                     4.3e-07;
                                                                          DB 22;
                                                                        Length 118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               augmenting
                                 <u>,</u>
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RESULT 3
AAB67513
                                        Region
                                                                           Key
Region
                                                                                                                                                                        autoimmune disease; infectious disease; inflammation; allergy; rheumatoid arthritis; myasthenia gravis; lupus erythematosus; multiple sclerosis; insulin-dependent diabetes mellitus; inflammation; transplant rejection; graft versus host disease.
                                                                                                                                                                                                                                                      Complementarity determining region; CDR; immune response; cytotoxic T lymphocyte associated antigen-4; CTLA-4; B7 1:
                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                   Heavy chain variable region of anti-CTLA-4 antibody 10D1
                                                                                                                                                                                                                                                                                                                                                                  29-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                          AAB67513;
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB67513 standard; peptide; 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50
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                                                                                                                                                                                                                                                                                                                                                               (first entry)
/note= "
99..107
                                                                           Location/Qualifiers
                                                        note=
                "CDR2"
                                                        "CDR1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   66
                                                                                                                                                                                                                                                        ligand; cancer;
                                                                                                                                                                                                                                                                             antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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THE XOX WAX WAX DXX AXX

(MEDA-) MEDAREX INC

01-MAR-2001. WO200114424-A2

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The present sequence represents the heavy chain variable region of CC human antibody 1001. This antibody specifically binds to human composition of the human antibody 1001. This antibody specifically binds to human composition of the human composition of the human composition of the antibodies are segonse to an antigen in a patient, where the antibodies block composition of human CTLA-4 to human 87 ligands. The antibodies are composition of the composition of T cells and for treating prostate composition of T cells and for treating prostate concer, melanoma or epithelial cancer. A polyvalent or polycional concer, melanoma or epithelial cancer. A polyvalent or polycional control preparation comprising two antibodies of the invention are useful for suppressing a immune response in a patient. They are used for treating cancer, infectious diseases and promoting beneficial autoimmune reactions for the treatment of diseases with inflammatory or allergic components. The polyvalent or polycional preparations are useful for treating autoimmune diseases such as rheumatoid arthritis, myssthenia gravis and lupus erythematosus, multiple sclerosis, insulin-dependent
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Best Local (
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lymphocyte associated antigen-4, useful for inducing, augmenting
prolonging immune response to antigen or for suppressing immune
                    25-MAR-1999;
                                                                   24-MAR-2000; 2000WO-US07946
                                                                                                                                                                                                                                                                                                                                 Human; neutralising antibody; interleukin-12; IL-12; antinflammatory; complementarity determining region; CDR; antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                     multiple sclerosis;
                                                                                                                                                                                                                                                                                          antisclerotic; neuroprotective; antiparasitic; antibacterial; if
                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-hIL12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB39830;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AB39830 standard; Peptide; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                 antibody heavy chain CDR2 amino acid sequence SEQ ID 346.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                    99US-0126603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transplant rejection,
                                                                                                                                                                                                                                                                     rheumatoid arthritis.
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Pred. No.
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                                                                                                                                                                                                                                                                                                               antipsoriatic; antiasthmatic; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and inflammation, graft versus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
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Antibody; heavy chain; CDR2;

bone resorption; loss of bone mass; bone tumour; obone cancer; rheumatoid arthritis; hypercalcaemia

osteoporosis.

osteoprotegrin binding protein; osteoclast format ion; loss of bone mass; bone tumour; osteoporosis;

complementarity

determining

formation; region; OPGbp; Antibody heavy chain variable region CDR2 #2.

17-DEC-2001

(first entry)

AAU08361;

AAU08361 standard; peptide;

17

23-FEB-2000;

2000US-0511139

23-FEB-2001; 2001WO-US05973.

30-AUG-2001.

WO200162932-A1

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RESULT 5
AAU08361
ID AAU0
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AC AAU0
XX
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                                                                                                                                                                                                                                                                                                                           CC This invention relates to a new human antibody specific for human CC interleukin-12 (IL-12). The invention also includes antigen binding portions that bind to IL-12. Sequences AAB39485-B35516 represent human CC anti-IL-12 antibody heavy and light chain complementarity determining CC region (CDR) amino acid sequences, and also includes variable region CC manino acid sequences. Other variable region amino acid sequences are CC given in AAB39517-B39560 and AAB40068-B40149. Sequences AAB39772-B40063 CC represent other CDR sequences. Light chain CDR3 consensus sequences are CC given in AAB40067. Primers used in the identification and CC construction of the antibodies of the invention are given in AAB40067. The antibody of the invention are given in AC61062-C61071. The antibody of the invention is a neutralising CC antibody and has antirheumatic; antiarthritic; antiaclerotic; CC cardiant; antiparasitic; neuroprotective; antipasoriatic; antiaclerotic; CC cardiant; antiparasitic; antipacterial and immunosuppressive activity. CT construction is disease, multiple sclerosis and relemantic darthritis. C especially Crohn's disease, multiple sclerosis and relemantic connection in the creatment of the antipation and the transmitter of a pharmacontic connection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Salfeld JG, Roguska M, Paskind M,
Kaymakcalan Z, Labkovsky B, Sakora:
Veldman GM, Venturini A, Warne NW,
Derbyshire EJ, Carmen S, Smith S,
                                                                                                                                                                                                                                                                                                                         Sequence
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GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                 also be used in the manufacture of a pharmaceutical composition
                                                                                                                                                                                                                                                                                                                                                               human IL-12 disorders.
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Y B, Sakorafas P, Priedrich S, Myles A,

Warne NW, Widom A, Elvin JG, Duncan AR,

S, Smith S, Holtet TL, Du Fou SL;
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Pred. No. 2.9e-07;
1; Mismatches 0
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                                                                                                                                                                                                                                            Antibody; OPGbp; human; P heavy chain; osteopathic; osteoprotegrin binding protein; osteoporosis; bone cancer; tumour; osteoporosis; bone cancer; rheumatoid arthritis; hypercalcaemia of malignancy; steroid-induced osteoporosis; mutant; mutein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an antibody or antigen binding domain (or tragment, variant or derivative), which binds to an osteoprotegrin binding protein (OPGbp) and which is an antagonistic antibody. The antibody or antigen binding domain may be administered to inhibit spacedast formation or activation, inhibit bone recorption in a mammal, rewent or treat loss of bone mass in a mammal and to prevent or treat consourcell growth in bone. The loss of bone mass results from osteoporosis, metastasis of cancer to bone, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                          Inti-OPGbp antibody heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypercalcaemia of malignancy and steroid-induced osteoporosis. The present sequence is an antibody heavy chain variable region complementarity determining region, CDR2, which can be incorporated an antibody/antigen binding domain of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibodies that bind antagonistically to osteoprotegrin binding protein, useful for treating osteoporosis, metastasis of cancer bone, rheumatoid arthritis, hypercalcaemia of malignancy and steroid-induced osteoporosis -
                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                /note= "Wild-type
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2..31
                                                                                                                                                                      Location/Qualifiers 2..117
                                           /note= "Wild-type
12
    note=
                                                                          note= "Wild-type Gln substituted
                                                                                                     note= "Framework region"
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chain P.
                                               The invention relates to an antibody or antigen binding domain (or fragment, variant or derivative), which binds to an osteoprotegrin binding protein (OPGbp) and which is an antagonistic antibody. The antibody or antigen binding domain may be administered to inhibit osteoclast formation or activation, inhibit bone resorption in a mammal prevent or treat loss of bone mass in a mammal and to prevent or treat loss of bone mass in a mammal and to prevent or treat tumour cell growth in bone. The loss of bone mass results from osteoporosis, metastasis of cancer to bone, rheumatoid arthritis, hypercalcaemia of malignancy and steroid-induced osteoporosis. The
 Sequence
                                                                                                                                                                                                              Antibodies that bind antagonistically to osteoprotegrin binding protein, useful for treating osteoporosis, metastasis of cancer bone, rheumatoid arthritis, hypercalcaemia of malignancy and
                                                                                                                                                                                                    protein, useful for treating osteoporosis, bone, rheumatoid arthritis, hypercalcaemia steroid-induced osteoporosis -
                                                                                                                                                                           Example 6; Fig 11; 239pp; English
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N-PSDB; AAS13367.
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22-FEB-2001; 2001US-0791153.
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            Published_Applications_AA:*

1: /cgn2_6/ptodata/1/pubpaa/US08

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US-10-194-975-24
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US-09-880-748-2055
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Sequence 1956, Ap
Sequence 982, App
Sequence 982, Appli
Sequence 7, Appli
Sequence 24, Appl
Sequence 24, Appl
Sequence 69, Appl
Sequence 69, Appl
Sequence 104, App
Sequence 1031, Ap
Sequence 2055, Appl
Sequence 1330, Ap
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Sequence 1935, Ap	1901	1887	e 10,	ce 24,	e 839,	e 983,	e 981,	e 977,	e 1731	e 1627	e 1519	e 1394	e 1201	e 1725	e 1724	e 1119	e 1117	e 1115	e 1113	e 1111	e 1110	e 1108	equence 1105	e 1102	Sequence 911, App

RESULT 1 US-09-880-748-1956

Sequence 1956, Application US/09880748 Publication No. US20030059937A1

GENERAL INFORMATION:

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; SOFTWARE: PatentIn Ver.; SEQ ID NO 1956; LENGTH: 249; TYPE: PRT; ORGANISM: Homo sapiens US-09-880-748-1956
                                                              RESULT 2
US-09-880-748-952
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Sequence 952, Application US/09880748
Publication No. US20030059937A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                           Query Match 89.2%;
Best Local Similarity 88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
                                                                                                                                                 50 FISYDGSHKYYADSVKG 66
                                                                                                                                                                                       1 FISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                         Score 83; DB 9;
Pred. No. 5.9e-05;
                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                               DB 9; Length 249;
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RESULT'4
US-10-078-958-7
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US-09-880-748-982
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Sequence 7, Application US/10078958
Publication No. US20030070185A1
                                                                                                                                                                                                                                                                                            SOPTWARE: PatentIn Ver.
SEQ ID NO 982
LENGTH: 251
                                                                                                                                                                            Best Local Similarity Matches 15; Conser
                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 89.2%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 952
                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/212,210 PRIOR FLING DATE: 2000-06-15 PRIOR APPLICATION NUMBER: 60/240,816 PRIOR FILING DATE: 2000-10-17 PRIOR APPLICATE: 2000-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-880-748-982
                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/293,499
                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/276,248
                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 3239
                                                                                                                                                                                                                                                                                   TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT: Ruben et al.
TILE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ruben et al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: PF523
TURRENT APPLICATION NUMBER: US/09/880,748
TURRENT FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ILE REFERENCE: PF523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    URRENT FILING DATE: 2001-06-15
RIOR APPLICATION NUMBER: 60/212,210
RIOR FILING DATE: 2000-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIOR APPLICATION NUMBER: 60/240,816 RIOR FILING DATE: 2000-10-17
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                                                                                                        50 FIRYDGSNKYYADSVKG 66
                                                                                                                                                                                                                                                                                                                                                                                                                 OR APPLICATION NUMBER: 60/277,379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
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                                                                                                                                        FISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09880748 o. US20030059937A1
                                                                                                                                                                            Conservative
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Pred. No. 5.9e-05;
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Pred. No.
                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                       Length 251
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RESULT 6 US-10-194-975-24

Sequence 24, Application US/10194975 Publication No. US20030039649A1 GENERAL INFORMATION:

APPLICANT: Foote, Jefferson
TITLE OF INVENTION: Super Humanized Antibodies
FILE REFERENCE: 501231.01

CURRENT APPLICATION NUMBER: US/10/194,975

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                                                                                                                          ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-194-975-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Variable amino acid US-10-078-958-7
                                                                Matches
                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                             SOFTWARE: PatentIn version 3.1 SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                              APPLICANT: FOOLE, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 08/759,620
PRIOR FILING DATE: 1996-12-03
NUMBER OF SEQ ID NOS: 79
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 7
LENGTH: 83
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                               Sequence 23, Ap
Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/194,975
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/305,111
PRIOR FILING DATE: 2001-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens FEATURE:
NAME/KEY: MOD_RES
LOCATION: (22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: PLURAL Vh AND Vk REG:
TITLE OF INVENTION: THEREFORM
FILE REFERENCE: CELL 4.18 CON
CURRENT APPLICATION NUMBER: US/10/078,958
CURRENT FILING DATE: 2002-02-19
                                                                                                                                                                               LENGTH: 98
                                                                                                                                                                                                                                                                                                              ITLE OF INVENTION: Super Humanized Antibodies
51 ISYDGSNKYYADSVKG 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 93.
nes 15; Conservative
                  2 ISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             36 ISYDGSNKYYADSVKG 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 ISYDGNNKYYADSVKG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INVENTION: TRANSGENIC INVENTION: PLURAL Vh INVENTION: THEREFROM
                                                                                                                                                                                                                                                                                                                                                                           Application US/10194975
No. US20030039649A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MENDEZ, MICHAEL J. GREEN, LARRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KUCHERLAPATI, RUKLAPHOLZ, SUSAN
                                                               Conservative
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                                                                            93.8%;
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                                                                            Score 82;
Pred. No. 3
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Pred. Nq. 2.9e-05;
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                                                                                          Length 98;
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Result
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Perfect score:
Sequence:
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Maximum DB seq length: 200000000
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Listing first 45 summaries
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           number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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Gapop 10.0 , Gapext 0.
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93
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                 US-08-211-202-118
US-08-211-202-111
US-08-211-202-141
US-08-31-3978-46
US-08-331-3978-46
US-08-331-3978-46
US-08-277-693-46
US-08-211-202-135
US-08-211-202-135
US-08-478-039-96
US-08-478-039-96
US-08-478-039-96
US-08-478-039-96
US-08-276-852-72
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US-08-276-852-77
US-08-276-852-77
US-08-899-575-74
US-08-899-575-74
US-08-899-575-74
US-08-899-575-75
US-08-899-575-75
US-08-899-575-77
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Sequence 118, App
Sequence 115, App
Sequence 146, Appl
Sequence 46, Appl
Sequence 46, Appl
Sequence 46, Appl
Sequence 315, App
Sequence 38, Appl
Sequence 38, Appl
Sequence 96, Appl
Sequence 97, Appl
Sequence 73, Appl
Sequence 74, Appl
Sequence 77, Appl
Sequence 77, Appl
Sequence 77, Appl
Sequence 77, Appl
Sequence 78, Appl
Sequence 79, Appl
Sequence 77, Appl
Sequence 77, Appl
Sequence 78, Appl
Sequence 79, Appl
Sequence 79, Appl
Sequence 71, Appl
Sequence 72, Appl
Sequence 73, Appl
Sequence 74, Appl
Sequence 75, Appl
Sequence 76, Appl
Sequence 77, Appl
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US-08-211-202-118
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ence 17,		Φ N	ο	e 139,	e 152,	16,	Sequence 77, Appl	76,	e 75,	74,	73,	72,	77,	76,	e 75,	10	Sequence 73, Appl

PRIOR APPLICATION DATA; APPLICATION NUMBER: GB 9206372.6 FILING DATE: 24-MAR-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: PCT/GB92/00883 FILING DATE: 15-MAY-1992 ATTORNEY/AGENT INFORMATION: NAME: DAY1d W. Clough REGISTRATION NUMBER: 36,107 REFERENCE/DOCKET NUMBER: 28111/31960 TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELEPHONE: 312-474-6300
PILING DATE: 23-SEP-1992 CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: GB 9120252.3 FILING DATE: 23-SEP-1991 PRIOR APPLICATION DATA: APPLICATION NUMBER: GB 9120377.8 FILING DATE: 25-SEP-1991 PRIOR APPLICATION DATA: APPLICATION NUMBER: GB 9206318.9 FILING DATE: 24-MAR-1992
Carifa a es.
TITLE OF INVENTION: PRODUCTION of chimeric antibodies - a TITLE OF INVENTION: combinatorial approach NUMBER OF SEQUENCES: 144 CORRESPONDENCE ADDRESS: ADDRESSEE: Borid W. Clough, Marshall O'Toole Gerstein Murray & ADDRESSEE: Borin STREET: 6300 Sears Tower, 233 South Wacker Drive CITY: Chicago
202-118 118, Application US/08211202 0. 5565332 INFORMATION: CANT: HOOGENBOOM, Hendricus Renerus Jacobus Matteus CANT: BAIER, Michael CANT: DESPERS, Laurent Stephane Anne Therese CANT: WINTER, Gregory Paul

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US-08-211-202-118
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                                                  REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                         FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                   PRIOR APPLICATION DATA:
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 ORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/211,202
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM FC
OPERATING SYSTEM:
                                        TELEPHONE:
                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: GB 9. STITING DATE: 23-SEP-1991
                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
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60606-6402
IER READABLE FORM:
Floppy disk
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                                                                                                                                     ICATION NUMBER
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                                                                                                                                                                                                                                                                                                                                                                           APPLICATION DATA:
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Illinois
                                                                       AGENI IN Clough
David W. Clough
NUMBER: 36,107
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amino acid
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WINTER, Gregory Paul
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NSTEM: PC-DOS/MS-DOS
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15-MAY-1992
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25-SEP-1991
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                                                   Query Match
Best Local Similarity
Matches 15; Conserv
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APPLICANT: Honjo, Tasuku
                                                                                                                                                                                                                               REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
                                                                                                                        MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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                   2 ISYDGNNKYYADSVKG 17
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US-08-331-398A-46 ; Sequence 46, Application US/08331398A : Patent No. 5608039

ENERAL INFORMATION:

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Result
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Gapop 10.0 ,
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A;Gene: GDB:IGKV3 A;Gene: GDB:IGKV3 A;Cross-references: GDB:136266 A;Map position: 2p12-2p11 C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (ka hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into l C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: autoantibody; chronic lymphocytic leukemia; heterotetramer; immunoglobulin C;Keywords: autoantibody; chronic lymphocytic leukemia; heterotetramer; immunoglobulin 밁 R;Goni, F.R.; Che J. Immunol. 142, Ig kappa chain V-III region (Pie) - human (fragment) C;Species: Homo sapiens (man) C;Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 21-Jan-2000 C;Accession: C30608 C;Species: Homo sapiens (man) C;Date: 30-Jun-1990 #sequence_revision 30-Jun-1990 #text_change 21-Jan-2000 Ś A; Molecule type: protein A; Residues: 1-108 < GON> A;Molecule type: mRNA A;Residues: 1-129 <KIP> A;Title: Structural and idiotypic characterization of the L chains A;Reference number: A30601; MUID:89215279; PMID:2496160 Ig kappa chain precursor V-III region (Hic) - human A;Status: preliminary A;Accession: C30608 Query Match Best Local S Matches 9 ;1-20/Domain: signal ;21-129/Product: Iq) Residues: 1-129 <KIP> Comment: The protein is Accession: PL0021 Superfamily: immunoglobulin V region; immunoglobulin homology; Keywords: heterotetramer; immunoglobulin Genetics: 90 1 QQYGSSPWT 9 Similarity Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson, 12, 3158-3163, 1989 Conservative 98 sequence E.; Chen, P.P.; Carson, D.A. 100.0%; one of chain the <u>.</u> Score 54; DB 2; Pred. No. 0.02; Mismatches (surface immunoglobulin M autoantibodies predicted nt chain variable region PMID:3127527 (H1C) <u>,</u> #status Length 108 Indels predicted <u>.</u> gene expressed in of human IgM autoanti Gaps expressed D.; Solc (kap

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C; species: Home sapiens (man)
C; : 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 20-Mar-1998
C; : 18-sion: S67940
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C,Superfamily: immunoglobulin V region; immunoglobulin homology
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                                                                                                                                                                                                                                                                                          ;Accession:
                                                                                                                                                                                                                                                                                                      ;Title: Cloning of a human ;Reference number: S67940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                . Exp. Med. 173, 461-470, 1991
flitle: Somatic mutations in the variable regions of a human
Reference number: JH0428, MUID:91108344; PMID:1899104
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24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 21-Jul-2000
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Pred. No. 0.61;
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Superfamily: immunoglobulin V region; immunoglobulin
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d to the EMBL Data Library,
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RESULT 2

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...munoglobulin V region; B
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SEQUENCE 108 AA; 11635
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***EDLINE=79221900; PubMed=111146;

**Seidman J.G., Max E.E., Leder P.;

**Pa kappa-immunoglobulin gene is formed by

**Without further somatic mutation.";

**Nature 280:370-375(1979).
                                                                                                                                                       MEDLINE=67056897; PubMed=4162931;
Gray W.R., Dreyer W.J., Hood L.;
"Mechanism of antibody synthesis: size
                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 1-33.
MEDLINE-77148916; PubMed=403522;
Burstein Y., Schechter I.;
"Anino acid sequence of the NH2-terminal extra piece segments of the precursors of mouse immunoglobulin lambda1-type and kappa-type light
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Primates;

    Last sequence update)
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    region B6.

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InterPro; IPR003006; Ig_MHC.
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Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
Immunoglobulin V region; Sigr
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InterPro; IPRO3596; Ig_v.
Pfam; PP00047; Ig; 1.
SMART; SM00406; IGv; 1.
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GLOBULIN ACTIVITY.
PIR; A01892; K3HUSI.
HSSP; P80362; 1WTL.
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"Amino acid sequence of the variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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P01620;
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Primates; Catarrhini;
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88.9%;
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Pred. No. 0.53;
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COMPLEMENTARITY-DETERMINING-1.
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COMPLEMENTARITY-DETERMINING-2.
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RESULT 5
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1: sp_archea:
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sp_virus:*
sp_vertebrate:*
sp_unclassified:*
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                                                          Q9j028 rabbit oral
056955 human papil
08x3x3 escherichia
Q9la41 shigella dy
Q9tin6 bacteriopha
Q9xjh9 bacteriopha
Q8z773 salmonella
                                                                                                                                                  Q9ul78 homo sapien
Q99y45 streptococc
Ol3664 schizosacch
O42996 schizosacch
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; 1g; 1.
SMART; SM00406; IGv; 1.
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HSSP; P80362; 1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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90 QOYGSSPLT 98
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109 AA;
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                                                                                                                                                                                                                                                           109
11646 MW;
                                                                                                                                                           75.9%;
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                                                                                                                        Score 41; DB 4; Length 109; Pred. No. 4:8; 0; Mismatches 1; Indels
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Last annotation update)
lin light chain variable
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                                                                                                                                                                                                                                                           5F675C52EC7EE197 CRC64;
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RESULT
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Best Local S
Matches 6
  Yeast 16:71-80(2000).

Yeast 16:71-80(2000).

EMBL, AB004539; BAA21455.1; ...

InterPro, IPR001680; WD40.

Pfam; PF00400; WD40; 1.

SMART; SM00320; WD40; 1.

PROSITE; PS00678; WD REPEATS 1; UNKNOWN 1.

PROSITE; PS50082; WD REPEATS 2; 1.

PROSITE; PS50082; WD REPEATS REGION; 1.
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Q99Y45;
01-JUN-2001
01-JUN-2001
                                                                                                                                                                                              Yanagida M., Noki K., Ogura K., F
Yanagida M.;
"A 38 kb segment containing th
Yeast chromosome II: sequence
genomic DNA and CDNAs encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JAN-1998 (TremBLrel.
01-JAN-1998 (TremBLrel.
01-JUN-2002 (TremBLrel.
kepeat;
                                                                                                                                                                                                                                                                                                                 Machida M., Yamazaki S.,
Haikawa Y., Yamazaki J.,
                                                                                                                                                                                                                                                                                                                                       MEDLINE=20089027;
Machida M., Yamaza
                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=972 H-;
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  phizosaccharomyces pombe (Fission yezst).

ukaryota; Fungi; Ascomycota; Schizosaccharomycetes;

Schizosaccharomycetales; Schizosaccharomycetaceae;

Schizosaccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pi073 protein.
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MEDLINE=21192684; PubMed=11296296;
Perretti J.J., McShan W.M., Ajdic D.J., Savic D.
Trimeaux C., Sezate S., Suvorov A.N., Kenton S.,
an Y., Jia H.G., Najar F.Z., Ren Q., Zhu H., &
uan X., Clifton S.W., Roe B.A., McLaughlin R.;
"Complete genome sequence of an M1 strain of Str.
Proc. Natl. Acad. Sci. U.S.A. 98:4658-4663 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Streptococcus pyogenes.
Streptococcus pyogenes.
Bacteria; Firmicutes; Bacillus/Clostridium group; Lactobacillales;
Streptococcaceae; Streptococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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InterPro; IPR000182; GCN5acetyltransf.
Pr00583; Acetyltransf; 1.
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Putative ribosomal-protein-alanine acetyltransferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CBI_TaxID=4896;
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Yamamoto s
K., Kudoh
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Last annotation update)
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Pred. No.
                                                                                                                                                                                           ne cdc2 gene from the left arm of analysis and characterization of on the segment.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an M1 strain of Streptococcus pyogenes.";
98:4658-4663(2001).
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                                                                                                                                                                                                                                                                                    Tanaka T., Ku
Sekine M., Og
Kikuchi H.,
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, Oguchi A., Nag
H., Zhang M.Q.,
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ai H.S., Lin
J L., White J
                                                                                                                                                                                                                                                                                                          Nagai Y.,
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RESULT 4
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ID 0429
AC 0429
AC 0429
DT 01-J
DT 01-
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Best Local S
Matches 6
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Best Local S
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Matches
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Pfam; PF00400; wD40; 1.

SMART; SM00320; wD40; 1.

PROSITE; PS00678; wD REPEATS 1; UNKNOWN 1.

PROSITE; PS50082; wD REPEATS 2; 1.

PROSITE; PS50294; WD REPEATS REGION; 1.

REPEAT; WD repeat.

SEQUENCE 391 AA; 44836 MW; A849BB001264D
SEQUENCE FROM N.A. Cla
Christensen N.D., Cla
Submitted (JAN-2000)
                                                                           SEQUENCE FROM N.A.
MEDILINE=20219637; PubMed=10753723;
Christensen N.D., Cladel N.M., Reed C
"Rabbit oral papillomavirus complete
following genital infection.";
Virology 269:451-461(2000).
                                                                                                                                                                                                                                                Papillomavirus.
                                                                                                                                                                                                                                                                                   Regulatory protein rabbit oral papillo
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                                                                                                                                                                                                                                                              rabbit oral papillomavirus
Viruses; dsDNA viruses, no
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                                                                                                                                                                                                                         CBI_TaxID=122291;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schizosaccharomyces pombe (Fission yeast).
Eukaryota, Fungi, Ascomycota, Schizosaccharomycetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WD repeat protein.
SPBC27B12.05.
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75.0%;
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                                                                                                                                                                                                                                                                RNA
                                                                                                                                                                                                                                                                                                                   Created)

Last sequence update)

Last annotation update)
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Pred. No. 18;
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Pred. No.
                                                                                                                                                                                                                                                            stage;
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1., Reed C.A., Han EMBL/GenBank/DDBJ

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Result
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Maximum Match 100%
Listing first 45 s
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Maximum DB seq length: 2000000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        humber of hits satisfying chosen parameters:
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                  /SIDS2/gcgdata/geneseq/geneseqp-embl/AA2001.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2002.DAT:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIDS2/gcgdata/geneseq/
                                                                                                                                                                                                 Length DB
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                                                      AAB67495
AAG93666
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AAR54307
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Human light chain
Human anti-Rh(D) a
Human HIV-1 monocl
Anti-HIV gp120 imm
VL region of HIV n
Anti-gp120 antibod
Anti-gp120 antibod
The kappa chain of
The kappa chain of
Light chain CDR3 f
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AAB62087	AAR30770	AAB06354	AAB06351	AAW24551	ū	AAP30251	AAR30880	ABP51796	AAG65862	AAW85910	AAW18271	AAR21310	AAW58478	AAW58505	AAW58515	AAR41283	AAR38614	ABB07233	ABB07231	AAU07490	AAU07472	AAR95306	AAR58533	AAY57337	AAY34039	AAW64671	5	AAW45517	æ	5	AAB62773	AAB62771	8	AAR95303
Ω	human	1 13	ΥĽ	#3 of r101-2	variable	e Off et	• '	prio	1-PrP anti	7 light	a	chai	anised an	ведие	Ø		modified	4.	-IL-4 recepto	antibo	Synthetic peptide	Light chain CDR3 o	Light chain CDR3 f	UC DANCA MAD NANUC	NANUC-2 antibody 1	Human UC pANCA mon	ative col	-2 light	tetanus tox	-Jk2. Homo	T-VIH	HIV-1 =	m light chai	Light chain CDR3 o

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RESULT 1
AAB67495
ID AAB6
XX
AC AAB6
XX
Complementarity determining region; CDR; immune response; antibody; cytotoxic Tlymphocyte associated antigen-4; CTLA-4; B7 ligand; cancer; autoimmune disease; infectious disease; inflammation; allergy;
                                                                                                                                                                                                                                                                                                                                AAB67495 standard; peptide; 9 AA.
                                                                         24-AUG-1999;
                                                                                             24-AUG-2000; 2000WO-US23356
                                                                                                                     01-MAR-2001.
                                                                                                                                        WO200114424-A2
                                                                                                                                                                                   multiple sclerosis; in
transplant rejection;
                                                                                                                                                                                                       rheumatoid arthriti
                                                                                                                                                                                                                                                             Human light chain complementarity determining region 3 (CDR3).
                                                                                                                                                                                                                                                                                     29-MAY-2001
                                                                                                                                                                                                                                                                                                           AAB67495;
                                                                                                                                                              Homo sapiens.
                                                  (MEDA-) MEDAREX INC.
                                                                                                                                                                                                          arthritie;
                                                                                                                                                                                                                                                                                    (first entry)
                                                                        99US-0150452
                                                                                                                                                                                             diabetes mellitus; inflammation;
                                                                                                                                                                                                          upus erythematosus
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WPI; 2001-202933/20 Korman AJ, Halk EL,

Lonberg N;

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RESULT
AAG936
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                     Human; anti-Rh(D) a
                                                                                                                                                 11-OCT-1996;
10-APR-1998;
27-JUN-1997;
 New \hat{k} solated protein, preferably a human diagnostics requiring a human instead of therapeutic medicine -
                                                           WPI; 2001-388931/41.
N-PSDB; AAH68723.
                                                                                                                                                                                                                                                                                                                                                                                                                                        JT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      an immune response to an antigen in a patient, where the antibodies also useful for treating autoimmune disease in a subject caused or exacerbated by increased activity of T cells and for treating prostate cancer, melanoma or epithelial cancer. A polyvalent or polyclonal antibody preparation comprising two antibodies of the invention are useful for suppressing a immune response in a patient. They are used for treating cancer, infectious diseases and promoting beneficial autoimmune mapponents. The polyvalent or polyclonal preparations for the treatment of diseases with inflammatory or allergic reating autoimmune diseases such as rheumatoid arthritis, myasthenia gravis and lupus erythematosus, multiple sclerosis, insulin-dependent diseases mellitus, transplant rejection, and inflammation, graft versus
                                                                                                 Siegel DL;
                                                                                                                                                                                                                             03-JUL-2001.
                                                                                                                                                                                                                                                      US6255455-B1
                                                                                                                                                                                                                                                                                                                                              Human anti-Rh(D)
                                                                                                                                                                                                                                                                                                                                                                       14-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                 AAG93666;
                                                                                                                          (UYPE-) UNIV
                                                                                                                                                                                                      29-JAN-1999;
                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                        AAG93666 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB67490-99 and AAB67501-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prolonging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human sequence antibody that binds to human cytotoxic lymphocyte associated antigen-4, useful for inducing, augmen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibodies are used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ш
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                                                                                                                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OOYGSSPWT 9
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune response to antigen or for suppressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                       PENNSYLVANIA
                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                                                96US-0028550.
98US-0081380.
97US-0884045.
                                                                                                                                                                                                  99US-0240274
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                                                                                                                                                                                                                                                                                                                                       antibody clone SH52 protein sequence
                                                                                                                                                                                                                                                                                                 antibody; Rh(D) binding protein; immunostimulant;
h phenotype; diagnosis; therapeutic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AB67501-06 represents complementarity determining human antibodies. The antibodies specifically billymphocyte associated antigen-4 (CTLA-4). Such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in methods for inducing,
                                                                                                                                                                                                                                                                                                                                                                                                                         108 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 54; DB 22;
Pred. No. 7.8e+05;
                 anti-Rh(D)
an animal a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 9;
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             antibody for use antibody and in
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Best Loc Matches

Query Match Best Local

Similarity

100.0%;

Conservative

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Score 54; DB Pred. No. 0.0 0; Mismatches

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Indels

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Gaps

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DB 22;

Length 108;

Sequence

108

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RESULT 3
AAB62770
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Best Local S
Matches 9
                The present invention provides the protein and coding sequences for the variable regions of human monoclonal antibodies which are immunoreactive with human immunodeficiency virus-1 (HIV-1) envelope glycoprotein gp120. These can be used in diagnosis and therapy of HIV-1 infection.
                                                                                                                            Novel human monoclonal antibody immunoreactive with human immunodeficiency virus-1 glycoprotein gp120, useful for detecting HIV-1 in biological sample and providing passive immunotherapy to HIV-1
                                                                                                                                                                                            WPI; 2001-112438/12.
N-PSDB; AAF29071.
                                                                                           Claim 1; Page 66; 81pp;
                                                                                                                          intected
                                                                                                                                                                                                                                       Watkins BA,
                                                                                                                                                                                                                                                                                               30-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus-1; HIV-1; human monocional antibody; envelope 91ycoprotein; 9p120; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preferably a human antibody, (I) having an amino acid sequence comprising one of the sequences (S) given in AAG9358 to AAG93669. (I) has immunostimulant activity, and can be used as an immune system stimulant. (I) can be used in diagnostic and therapeutic medicine. The antibodies used in diagnostics that require human antibodies instead of animal antibodies, such as determine the Rh phenotype of human red blood cells. AAHG8615 to AAHG9726 represent the nucleotide sequence which encode AAG93558 to AAG93669. AAG93670 to AAG93697 represent anti-Rh(D) heavy than CDR3 amino acid sequences which are given in the exemplification
                                                                                                                                                                                                                                                             (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                          23-JUN-2000; 2000WO-US17327.
                                                                                                                                                                                                                                                                                                                                                         04-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB62770;
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                                                                                                                                                                                                                                                                                                                                                                                  WO200100678-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              68
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                                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QQYGSSPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QQYGSSPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
9; Conserv
                                                                                                                          mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; Protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monoclonal antibody
                                                                                                                                                                                                                                       Reitz MS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
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                                                                                                                                                                                                                                                                                             99US-0141701.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      70; 162pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          describes an isolated Rh(D) binding protein,
                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 54;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 108;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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문 Ś

88

QQYGTSPWT QQYGSSPWT

96 9

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RESULT 4
AAR54307
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      Best Local Similarity
Matches 8; Conserv
                                            Query Match
                                                                                                                                  Lymphocyte mRNA was converted to cDNA and subjected to PCR amplification using primers specific for heavy and light chain variable regions. The amplification products were inserted into a dicistronic vector to produce a library of fragments. E.coli Xii Blue cells were transformed with the library. Filamentous phage were produced which expressed the MAb regions on their surface. Panning with gpl20 and gp41 resulted in the recovery of immunoreactive clones. The light chain VK region sequence AAR54307 is from a gp120-
                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                    New human monoclonal antibodies neutralising HIV - react with gp120 or gp41 and nucleic acid encoding them, useful for in vivo or in vitro diagnosis and for passive immuno-therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus; HIV1; glycoprotein gp120; epitope neutralisation; monoclonal antibody; kappa light chain; variable region; framework; complementarity determining region.
                                                                                                                                                                                                                                                                                                                                Example; Page 177-178;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-APR-1994
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                                                                                                                          specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..PI; 1994-135516/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR54307 standard; protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           omo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OQYGSSPWT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QQYGSSPWT
                                                                                                                            clone.
                                                                                    108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Burton DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-0954148
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56..87
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                       94.4%;
                                                                                                                                                                                                                                                                                                                            248pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lerner
Score 51; DB 15; Length 108; Pred. No. 0.26; Indels
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Conservative

٠, Gaps

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Query Match
Best Local Similarity
Matches 8; Conserv

94.4%; llarity 88.9%; Conservative

Score 51; DB 17; Pred. No. 0.26; 1; Mismatches 0

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Gaps

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Length 108; Indels

Sequence

108

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                      The sequences given in AAW01261-92 represent the light chain variable regions (VL) of a series of monoclonal antibodies (MAb's) which are immunoreactive with HIV glycoprotein gp120 and are capable of neutralising HIV. This sequence represents the sequence of the JK1 gene clone, b24. A MAb containing this VL sequence has the capacity to reduce HIV infectivity titre in an in vivo virus infectivity assay by 50 % at a concentration of less than 700 mg of antibody(ml, and binds mature gp120 preferentially over the precursor gp160. The MAb may be used for determining immunocompetence of a human anti-HIV antibody and in the detection of HIV infection.
                                                                                                                                                                                                       Monoclonal antibody binding to V1/V2 loop of HIV gp120 -passive immuno:therapy and detection of HIV infection.
                                                                                                                                                                              Example; Fig 11; 366pp; English
                                                                                                                                                                                                                                                      WPI; 1996-179601/18
                                                                                                                                                                                                                                                                                    Barbas CF,
                                                                                                                                                                                                                                                                                                                                          18-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                  01-FEB-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heavy chain; light chain; variable region; VH; monoclonal antibody; MAb; HIV; human immunodeficiency virus; glycoprotein; gpl20; clone; virus infectivity assay; precursor gpl60; immunocompetence; human; anti-HIV antibody; detection; HIV infection.
                                                                                                                                                                                                                                                                                                                                                                    11-JUL-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                              (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VL region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW01265 standard; Protein; 108 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of HIV neutralising MAb,
                                                                                                                                                                                                                                                                                 Burton DR,
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                                                                                                                                                                                                                                                                                                                                         94US-0276852
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        97..108
/label=
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                                                                                                                                                                                                                                                                                 Lerner RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FR4
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밁 S

88

QQYGTSPWT OOYGSSPWT

96

Anti-gp120 antibody light chain variable

region from

clone

b24

30-JUN-2000 AAY95117

(first entry)

AAY95117 standard; Protein; 108 AA.

Ş 뭐

QQYGSSPWT

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cc a human whole immunoglobulin (Ig) molecule which immunoreacts with HIV compatible immunoglobulin (Ig) molecule which immunoreacts with HIV cc gp160 and neutralises HIV and which reduces HIV infectivity titre in an cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity assay by 50%, at a concentration of less than cc in vitro virus infectivity as the diagnosts and cc infection in vitro virus infectivity and infectivity in the cc inmunocharal antibodies are useful information regarding the cc which can be used to screen human monoclonal antibodies to identify composition. The neutralising antibodies define new epitopes on the invention. The neutralising antibodies define new epitopes on the immunotherapeutic human monoclonal antibodies define new epitopes on the cc immunotherapeutic human monoclonal antibodies. A major advantages of the monoclonal antibodies for diagnosis and immunotherapy of HIV induced disease greatly composition and infective in vivo use of the monoclonal cc antibodies for diagnosis and immunotherapy of HIV induced disease greatly administered antibodies which is a problem commonly encountered when cc antibodies for diagnosis and immunotherapy of the monoclonal antibodies described derives from the fact that they inmunoreact with a unique determinant cc present on mature HIV glycoprotein gp120. This class of antibodies is a particularly effective at neutralising field isolates of HIV.
Query Match
Best Local Similarity
-- has 9, Conserve
                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; Figure 11; 366pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nove1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-human immunodeficiency virus type 1 antibody; HIV-1; neutralise; reduce HIV infection; diagnosis; immunotherapy; HIV induced disease; glycoprotein 120; gp120; glycoprotein 41; gp41; monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU9948756-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .mmunodeficiency virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human monoclonal antibodies which immunoreact with and neutralise immunodeficiency virus useful for treating HIV infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DR,
                                                                                                   108
                 94.4%;
ilarity 88.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence represents a fragment of an anti-human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99AU-0048756
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            Score 51; DB
Pred. No. 0.26
1; Mismatches
                                                   DB 21;
                                                   Length 108;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates
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RESULT 7
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ID AAY9
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XX AAY9
XX AAT1
XX ITMM
XX AAT1
XX AAT1
XX AAT1
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XX ITMM
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XX A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antibody; anti-HIV monoclonal antibody; glycoprotein-120; human immunodeficiency virus type 1; HIV-1; infectivity t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-gp120 antibody light chain variable region from clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       passive immunotherapy; reduce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mmunocompetence; active immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Barbas CF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lerner RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    severity; HIV-induced
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Human neutralizing monoclonal antibodies to human immunodeficiency virus (HIV) used for providing passive immunotherapy to HIV are specific for glycoprotein-120 -

Example 9; Figure 11; 374pp; English.

Immunotherapy to HIV in a human. They can be administered to high-risk patients to reduce the likelihood and/or severity of HIV-induced disease of and to patients who are already HIV-infected. The antibodies are used to neutralising field isolates which provides information about the immunocompetence of an immune response in HIV patients, for detecting producing anti-idiotypic antibodies which can be used for active immunosation and to screen human monoclonal antibodies to identify those with the same binding specificity and to monitor the course of HIV present in the body or in body fluids by immunoassay. The anti-HIV gp-120 connoclonal antibodies are used to in the body or in body fluids by immunoassay. The anti-HIV gp-120 connoclonal antibodies are encoded by a human polynucleotide sequence and The invention relates to the production of an anti-HIV (human immunodefictency virus) glycoprotein (gp)-120 monoclonal antibody capable of reducing an HIV infectivity titre in an in vitro virus infectivity assay by 50% at a concentration of less than 70 ng/ml. method for the production of the antibody comprises:

(a) providing a first polymucleotide encoding a heavy chain immunoglobulin amino acid sequence (which does not comprise the section temperated by AAY98206) and a second polymucleotide encoding a light immunoglobulin amino acid sequence (which does not comprise the section immunoglobulin amino acid sequence) sequences cell; and This sequence represents a fragment of the The invention relates to the production of (d) isolating (c) maintaining the host cell in conditions which allow the amino acid in immunoglobulin amino acid sequence, inserting the first and second polynu the antibody comprising the heavy and light chain and second polynucleotide sequences into a host antibodies of the invention sequence light The

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Maximum Match 100%
Listing first 45 summaries
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               April 16, 2003, 17:21:54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB_pep:*
/cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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6/ptodata/1/pubpaa/PCT NEW PUB.pep:*
6/ptodata/1/pubpaa/US06_NEW PUB.pep:*
6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
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                                                                                US-09-848-798-178
US-09-056-593-118
US-09-056-160B-12
US-09-056-160B-12
US-09-811-123-6
US-10-237-667-18
US-10-237-876-18
US-10-237-876-18
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US-09-782-397-5
US-09-782-397-14
US-09-782-397-14
US-09-782-397-14
US-09-782-397-14
US-09-782-397-14
US-09-828-708-52
US-09-828-708-52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Search time 2.65263 Seconds
(without alignments)
256.547 Million cell updates/sec
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Sequence 178, App
Sequence 118, App
Sequence 74, Appl
Sequence 12, Appl
Sequence 6, Appli
Sequence 18, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 175, Appl
Sequence 125, Appl
Sequence 52, Appl
Sequence 54, Appl
Sequence 55, Appl
Sequence 56, Appl
Sequence 7, Appl
Sequence 7, Appl
Sequence 7, Appl
Sequence 10, Appl
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39	39	39	39	39	39	39	39	39	39	39	39	39	9	39	39	υ 9	ω 9	9	39	39	39	39	40	40	40	
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100,	e 1226	1226,	117,	115,	113,	111,	Sequence 109, App	107,	105,	103,	126,	10,	ø	Sequence 16, Appl	4	Sequence 2, Appli	16,	Sequence 4, Appli	Sequence 2, Appli	5,	13,	e 6,		99	Sequence 12, Appl	

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RESULT 2
US-10-006-593-118
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                                                                                                                                                                                                                                                                                                                                                                                               ; FEATURE:
; OTHER INFORMATION: anti-Rh(D) antibody clone SH52
US-09-848-798-178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/848,798

CURRENT FILING DATE: 2001-05-04

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274

PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29

PRIOR PRIOR ON NUMBER: EARLIER APPLICATION NUMBER: 60/028,550

PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11

NUMBER: OF SEQ ID NOS: 224

SOCTWARE: PATENTIN Ver. 2.0

SEQ ID NO 178

LENGTH: 108
                                                                 Sequence 118, Application US/10006593 Publication No. US20030049683A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 09596-42U2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 178, Application US/09848798 Publication No. US20030040605A1 GENERAL INFORMATION:
APPLICANT: Bowdish, Katherine S
APPLICANT: Frederickson, Shana
APPLICANT: Renshaw, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                      68
                                                                                                                                                                                                                                                   1 QQYGSSPWT 9
                                                                                                                                                                                                                      QQYGSSPWT 97
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Donald L.
                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                     Score 54; DB 9;
Pred. No. 0.022;
Mismatches
                                                                                                                                                                                                                                                                                                                                              DB 9; Length 108;
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; ORGANISM: human
US-10-006-593-118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TLE OF INVENTION: RATIONALLY DESIGNED ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 QQYGSSPW 8
MÓSECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 74:
                                                                                                                                TELEX: <Unknown>
WATION FOR SEQ ID NO: 74:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                  REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06510/059001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/288,889
FILING DATE: 2001-05-04
APPLICATION NUMBER: US 60/294,068
FILING DATE: 2001-05-29
                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Prusiner, Stanley B. Williamson, R. Anthony
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PatentIn version 3.1
                                                                                    LENGTH: 109 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 2.0
                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PLICATION NUMBER: US/10/006,593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
                                                                                                                                                                                                ELEPAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Fish & Richardson P.C. STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/09943906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATION NUMBER: US 60/251,448
DATE: 2000-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burton, Dennis R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-12-05
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RESULT 5 US-09-811-123-6

Sequence 6, Application US/09811123
Patent No. US20020001587A1

INFORMATION:

NEKAL INFORMATION ETICKBON
IPPLICANT: Sharon ErickBon
IPPLICANT: Ralph Schwall
APPLICANT: Mark Sliwkowski
PTTLE OF INVENTION: METHODS OF TREATMENT USING ANTI-EYDB

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US-09-056-160B-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
                                                     Query Match
Best Local Similarity 77.1
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 79.6%;
Best Local Similarity 87.5%;
Matches 7; Conservative
                                                                                                                                                                                     TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/0 FILING DATE: 06-AUG-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
89 QQYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORRESPONDENCE ADDRESS
                                                                                                                                         LENGTH: 108 amino acids
                              1 QQYGSSPWT 9
                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                           REGISTRATION NUMBER: 28
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                               FELEPHONE:
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/056,160B
FILING DATE: 06-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 OYGSSPWT 9
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                                                                                                                             Linear
                                                                                                                                                                                                                                                                                                                                                                                                                              IBM PC compatibl
                                                                                                                                                                                                              650/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Presta, Leonard G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0032315A1
                                                                                                                                                                                                                                                                                                                                                          06-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yvonne M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTI-VEGF ANTIBODIES
                                                                   77.8%;
77.8%;
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                                                                                                                                                                                                                                                      28,616
                                                                                                                                                                                     12:
                                                                                                                                                                                                                                                                                                                                                                                                                                          1.44 Mb floppy disk
                                                         0
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                                                                     Score 42; DB 10; Length 108; Pred. No. 2.6;
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Pred. No.
                                                        Mismatches
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Title:
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Sequence:
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Maximum Match 100%
Listing first 45 s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Database
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                22643211111112
226432118
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Gapop 10.0 , Gapext 0.5
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US-08-350-260A-503
US-08-329-525-26
US-08-480-753-6
US-09-041-889-11
US-08-301-645-50
PCT-US94-01258-50
PCT-US94-01258-50
US-08-488-1123-123
US-08-488-113B-123
US-08-488-113B-125
US-08-488-113B-125
US-08-477-484B-123
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US-08-17-669D-27
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US-08-276-852-86
US-08-899-575-86
US-08-899-575-86
PCT-US95-08743-86
US-08-652-5584-9
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Sequence 86, 1
Sequence 86, 1
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Patent No. 62
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; OTHER INFORMATION: anti-Rh(D) antibody clone US-09-240-274-178
                                                                                                                  GENERAL INFORMATION:
APPLICANT: Barbas, C
APPLICANT: Lerner, R
APPLICANT: Lerner, R
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. SEQ ID NO 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 9; Conser
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CURRENT FILING DATE: 1999-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF FILE REFERENCE: 09596-42U2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                                                              CORRESPONDENCE ADDRESS
                                                                                                 NUMBER OF SEQUENCES:
                                      ADDRESSEE: Patent Counsel
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10666 No.
Mail Drop
                                                                                                                                                                                                                                                               Application US/08276852
                                                                                               Burton, Dennis R
Barbas, Carlos F
Lerner, Richard A
ZENTION: RICHARD NEUTRALIZING MONOCLONAL ANTIBODIES
ZENTION: TO HUMAN IMMUNODEFICIENCY VIRUS
QUENCES: 170
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                                                       The Scripps Research Institute, Office of
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5652138th Torrey Pines Road, Suite 220, TPC8
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COUNTRY:

La Jolla ·CA Y: USA

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Best Local Similarity
Marches 8; Conserve
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                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURREINT APPLICATION DATA:
                                                                                                                 COMPUTER READABLE FORM:
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APPLICATION NUMBER: US 07/954,148
FILING DATE: 30-SEP-1992
TTORNEY/AGENT INFORMATION:
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TYPE: amino acid
TOPOLOGY: linear
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OPERATING SYSTEM:
                                                                                                 EDIUM TYPE:
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                                                                                                                                  COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 0 PILING DATE: 30-SEP-1993
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FILING DATE: 18-JUL-1994
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                 CATION NUMBER:
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10666 No. 5770440th Torrey Pines Road, Suite 220,
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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24-JUL-1997
                                                                                                                                                                                                                                                                                       Richard A
HUWAN NEUTRALIZING MONOCLONAL ANTIBODIES
TO HUWAN IMMUNODEFICIENCY VIRUS
             US/08/899,575
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tent No.
                                 APPLICATION NUMBER: US 07/954,148 FILING DATE: 30-SEP-1992 ATTORNEY/AGENT INFORMATION:
                                                                               APPLICATION NUMBER: US 08/276,852
FILING DATE: 18-UU1-1994
APPLICATION NUMBER: US 08/178,302
FILING DATE: 30-SEP-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/899,575
FILING DATE: 24-JUL-1997
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REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                            CLASSIFICATION:
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TLE OF INVENTION:
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TOPOLOGY: linear
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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58
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1: pir1:*
2: pir2:*
3: pir3:*
Minimum Match 0%
Maximum Match 100%
Listing first 45 so
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AF2650
A82366
AH0482
G65146
D91174
D86020
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S33553
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RESULT
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C;Species: Mus musculus (house mouse)
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 05-Nov-1999
C;Accession: I56197; I49661
                                                                                                                                                                                    C; Keywords: immunoglobulin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Ahouse, J.J.; Hagerman, C.L.; Mittal, P.; Gilbert, D.J.; Copeland,
                                                                                                                                                                                Gene: FCRn superfamily: class I histocompatibility antigen; immunoglobulin homology
                                                                                                                                                                                                                                                                                                       ;Molecule type: mRNA;Residues: 1-72,'M',
                                                                                                                                                                                                                                                                                                                                                                                        ;Title: Structural and phylogenetic analysis of the MHC class I-like Fc receptor gene.;Reference number: I49661; MUID:95270984; PMID:7538537
                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Cross-references: GB:L17022; NID:g293385; PIDN:AAA16904.1; PID:g293386; Kandil, B.; Noguchi, M.; Ishibashi, T.; Kasahara, M. Immunol. 154, 5907-5918, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . Immunol. 151, 6076-6088, 1993 ;Title: Mouse MHC class I-like Fc receptor encoded outside the MHC ;Reference number: I56197; MUID:94065162; PMID:7504013
                                                                                                                                                                                                                                                                     Cross-references: GB:D37874; NID:g1009353; PIDN:BAA07111.1; PID:g1009354
                                                                                                                                                                                                                                                                                                                                                  Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local :
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  48
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TGWLGPQQY 56
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77.8%;
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Pred. No. 5
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A;Reference number: A64200; MUID:96026346; PMID:7589993
A;Accession: F64230

preliminary; nucleic acid sequence not shown; translation

not shown

spore germination apparatus protein (gerBB) homolog - Mycoplasma genitalium C;Species: Mycoplasma genitalium C;Date: 17-Nov-1995 #sequence_revision 17-Nov-1995 #text_change 21-Jul-2000 C;Accession: F64230

R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleischmann, R.D.; M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; Phillips, C.A.; Merrick, J.;

, C.A.; Venter, J.C. Science 270, 397-403, 1995 A;Title: The minimal gene

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Karp, F.; Science 294,
                                                                                                                                                                                                                                                                                                                       C;Accession: AE3245
R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks
erage, G.; Gillet, W.; Grant, C.; Guenthner,
                                                                                                     A;Gene: accE
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A;Start codon: GTG
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A;Residues: 1-970 <TIGR>
A;Cross-references: GB:U39707; GB:L43967; NID:g3844865; PIDN:AAC71499.1; PID:g1045972;
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                                                                      Genome: plasmid;SuperfamAly: ol
          Matches
                     Query Match
Best Local Similarity
                                                                                                                                  Experimental source: strain
                                                                                                                                             Cross-references: GB:AE008690; PIDN:AAL46379.1;
                                                                                                                                                                                                 Status: preliminary
                                                                                                                                                                                                                :Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58:Reference number: AB2577; PMID:11743193
Accession: AE3245
                                                                                                                                                                                                                                                                              Authors: Yoo, H.; Tao,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily: oligopeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;Residues: 1-286 <PIP>;Cross-references: EMBL;AF010180; NID:g3153171; PIDN:AAC17201.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Piper, K.R., Beck von Bodman, S., ubmitted to the EMBL Data Library,
                                                                                                                                                                     Residues:
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Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 08-Oct-199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              932
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85.7%;
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Pred. No.
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Pred. No. 23;
1; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                  R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cook, D.M.; Hwang,
May 1998
 Mismatches
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RESULT 7
JC5604
ABC-transporting peroxisomal membrane protein 69 - human
C;Species: Homo sapiens (man)
C;Date: 23-Sep-1997 #sequence_revision 23-Sep-1997 #text_change 02-Feb-2001
C;Accession: JC5604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;DelVecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucel.
                                                                                                                                                                                                                                                                                                                                              A;Gene: BMEII0129
A;Map position: I
                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-214 <KUR>
A;Residues: 1-214 <KUR>
A;Cross-references: GB:AE008918; PIDN:AAL53370.1; PID:g17984261; GSPDB:GN00191
A;Experimental source: strain 16M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Reference number: AD3252; PMID:11756688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hydrolase family protein BMEII0129 [imported] - Brucella melitensis (strain
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F;WOLff, G.; Plante, I.; Lang, B.F.; Kueck, U.; Burger, G.
J. Mol. Biol. 237, 75-86, 1994
A;Title: Complete sequence of the mitochondrial DNA of the
A;Reference number: Z17373; MUID:94180393; PMID:8133522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NADH2 dehydrogenase (ubiquinone)
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Species: Brucella melitensis
Date: 01-Feb-2002 #sequence :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;Superfamily: NADH dehydrogenase (ubiquinone) chain 1;Reywords: membrane-associated complex; mitochondrion; NAD; oxidative phosphorylatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Status: preliminary
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| Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change
                                                                                                                                                                                                                                                                                                                                            position:
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                                                                                                                                                                                                                                                            Conservative
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85.7%;
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HMAA_TRICA
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EMBL; D37874; BAA07111.1; EMBL; D37873; BAA07110.1; EMBL; D37872; BAA07110.1;

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Q49409; Q49253;
Q1-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
16-CCT-2011 Rel. 40, MG277.
                                                 SEQUENCE OF 624-678 FROM N.A.
STRAIN-ATCC 33530 / G-37;
MEDLINE-94075230; PubMed-8253680;
Peterson S.N., Hu P.-C., Bott K.F., Hut
"A survey of the Mycoplasma genitalium
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Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycoplasma.
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MGD; MGI:10301
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 Bacteriol. 175:7918-7930(1993).
SUBCELLULAR LOCATION: Integral
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EXTRACELLULAR ALPHA-2.
EXTRACELLULAR ALPHA-3.
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                          J. Biol. Chem. 272:22979-22989(1997).

-I FUNCTION: VISUAL PIGMENTS ARE THE LIGHT-ABSORBING MOLECULES THAT MEDIATR VISION. THEY CONSIST OF AN APOPROTEIN, OPSIN, COVALENTLY LINKED TO CIS-RETINAL.

-I SUBCELLULAR LOCATION: Integral membrane protein.
-I- TISSUB SPECIFICITY: RETINA. EXPRESSED IN THE DEPOLARIZING CELL LAYER OF THE PHOTORECPTOR CELLS DISTANT FROM THE LENS.
-I- PIM: SOME OR ALL OF THE CARBOXYL-TERMINAL SER OR THR RESIDUES MAY BE PHOSPHORYLATED (BY SIMILARITY).

-I- SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                                                                                                                MEDLINE=97435252; Pu
Kojima D., Terakita
Shichida Y.;
                                                                                                                                                                                                                                                                                                        OPS1_PATYE
O15973;
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DOMAIN
CONFLICT
SEQUENCE
                                                                                                                                cells.
                                                                                                                                                                                  TISSUE=Eye;
                                                                                                                                                                                                                   Patinopecten yessoensis (Ezo giant scallop) (Yesso scallop).
Bukaryota; Metazoa; Mollusca; Bivalvia; Pteriomorphia; Pectinoida;
Pectinoidea; Pectinidae; Mizuhopecten.
                                                                                                                                                                                                                                                                                                                             PATYE
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5-JUL-1998 (Rel.
0-MAY-2000 (Rel.
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ta A., Ishikawa
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SWISS-PROT entry is copyright.

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sp_unclassified:*
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Q8U604
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Q37629
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PRINTS; PR01638; MHCCLASSI.
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EMBL; D37908; BAA07137.1; JOINED.
HSSP; P13599; 3FRU.
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Kandil E., Noguchi M., Ishibashi T., F
"Structural and phylogenetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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Mus musculus (Mouse).
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25 TGWLGPQQY 33
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177 AA;
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Rodentia;
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MEDITE-95113783; PubMed=7814335;
Hwang I., Cook D.M., Farrand S.K.;
"A new regulatory element modulates homoserine
autoinduction of Ti plasmid conjugal transfer."
J. Bacteriol. 177:449-458(1995)
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01-DEC-2001
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Plasmid pTiC58.
Bacteria, Proteobacteria,
Rhizobiaceae, Rhizobium.
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NCBI_TaxID=176299;
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Mammalia; Eutheria;
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8; MHCCLASSI.
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Sciurognathi; Muridae;
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MEDLINE=98193120; PubMed=9524202;
Suzuki K., Ohta N., Hattori Y., Uraji M., Katoh A.,
"Novel structural difference between nopaline- and of
gene:construction of genetic and physical map and so
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE;
Plasmid.
                                                                                    SEQUENCE FROM N.A. STRAIN=MAFF301001;
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STRAIN=MARF301001;
MEDLINE=20184752; PubMed=10721727;
Suzuki K., Hattori Y., Uraji M., O
Katoh A., Yoshida K.;
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01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
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SEQUENCE FR
STRAIN=C58;
                                                                                                                                                                                                                                                                                                                                   Agrobacterium tumefaciens.
Plasmid pTi-SAKURA.
Bacteria; Proteobacteria;
Baizobiaceae; Rhizobium.
                                                                                                                                                  Gene 242:331-336(2000).
                                                                                                                                                                   "Complete nucleotide sequence of a
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InterPro, IPR000515; BPD_transp; 1.

Pfam; PF00528; BPD_transp; 1.

PROSITE; PS00402; BPD_TRANSP_INN_MEMBR;
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"Characterization of the acc operor pTiC58, which encodes utilization ( susceptibility to agrocin 84.";
J. Bacteriol. 179:7559-7572(1997).
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"The tra region of the nopaline-type Ti plasmid is a chimera with
elements related to the transfer systems of RSF1010, RP4, and F.";
J. Bacteriol. 178:4233-4247(1996).
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Listing first 45 summaries
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Match Length
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Gapop 10.0 , Gapext 0.5
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58
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The heavy chain of
Peptide #1263 enco
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KW cytotoxic T lymphocyte associated antigen 4; CTLA-4; BT ligand; cancer; KW autoimmune disease; infectious disease; inflammation; allergy; KW rheumatoid arthritis; mysathenia gravis; lugus erythematosus; KW multiple sclerosis; insulin-dependent diabetes mellitus; inflammation; XX transplant rejection; graft versus host disease. XX WO200114424-A2. XX PD 01-MAR-2001. XX PF 24-AUG-2000; 2000WO-US23356. XX PR 24-AUG-1999; 99US-0150452. XX PR 24-AUG-1999; 99US-0150452. XX PR (MEDA-) MEDAREX INC. XX Korman AJ, Halk EL, Lonberg N;							* *	•					DE Human heavy chain complementarity determining region 3 (CDR3)	DT 29-MAY-2001 (first entry)	AC AAB67492;	YY AAB67492 standard; peptide; 9 AA.	AAB67492	RESULT 1	-
							180.	s mellitus; inflammation:	is erythematosus:	ummation: allergy:	CTLA-4; B7 ligand; cancer:	me response; antibody;	region 3 (CDR3)						

39.5 39.5

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AAB67514
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Best Local
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                                                     Key
Región
                                                                                                                                                         Complementarity determining region; CDR; immune response; antibody; cytotoxic Tlymphocyte associated antigen-4; CTLA-4; B7 ligand; cancer; autoimmune disease; infectious disease; inflammation; allergy; rheumatoid arthritis; myasthenia grav4s; lupus erythematosus; multiple sclerosis; insulin-dependent diabetes mellitus; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer, melanoma or epithelial cancer. A polyvalent or polyclonal antibody preparation comprising two antibodies of the invention are useful for suppressing a immune response in a patient. They are used for treating cancer, infectious diseases and promoting beneficial autoimmune reactions for the treatment of diseases with inflammatory or allergic components. The polyvalent or polyclonal preparations are useful for treating autoimmune diseases such as rheumatoid arthritis, myasthenia gravis and lupus erythematosus, multiple sclerosis, insulin-dependent diabetes mellitus, 'transplant rejection, and inflammation, graft versus hong disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is present sequence represents the heavy chain variable region of sman antibody 1001. This antibody specifically binds to human crystotoxic T lymphocyte associated antigen-4 (CTLA-4). Such antibodies are used in methods for inducing, augmenting or prolonging an immune binding of human CTLA-4 to human B Tigands. The antibodies are also useful for treating autoimmune disease in a subject caused or exacerbated by increased activity of T cells and for treating prostate cancer, melanoma or epithelial cancer. A polyvalent or polyclonal
              Region
                                                                                                                                                                                                                                                                                Heavy chain variable region of anti-CTLA-4 antibody 10D1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                             29-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                      AAB67514;
                                                                                                                                                                                                                                                                                                                                                                                                      AAB67514 standard; peptide; 118
                                                                                                                                               transplant rejection; graft
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human sequence antibody that binds to human cytotoxic lymphocyte associated antigen-4, useful for inducing, augmen prolonging immune response to antigen or for suppressing imm response in patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGWLGPPDY 107
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ilarity 100.0%;
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              /note= "CDR1"
50..66
                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lomberg
                                                                                                                                               versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 58; DB 22;
Pred. No. 0.014;
Mismatches 0;
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RESULT 7
AAY72375
ID AAY7
XX AAY7
AC AAY7
XX
DT 24-A
XX
DE Amph
XX
Reco
KW Stab
KW COSH
XX
OSH
XX
Synt
XX
PD E910
XX
PD 27-1

Amphiphilic

24-APR-2001 AAY72375;

(first entry)

AAY72375 standard; Protein; 599 AA.

Recombinant protein; collagen-like polymer; gelatin-like polymer; NIN2P4; stabiliser; oil-in-water emulsion; foodstuff; pharmaceutical product;

recombinant collagen-like polymer, N1N2P4.

cosmetic product; photography

27-DEC-2000 EP1063565-A1 Synthetic. 밁

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                                                                                                                                                The present sequence represents the heavy chain variable region of CC human antibody 1001. This antibody specifically binds to human CY that it is a specifically binds to human CY that it is a specifically binds to human CY that it is a sugmenting or prolonging an immune CY response to an antigen in a patient, where the antibodies block CC binding of human CYLA-4 to human B7 ligands. The antibodies are CY also useful for treating autoimmune disease in a subject caused or cancer, melanoma or epithelial cancer. A polyvalent or polyclonal CY cancer, melanoma or epithelial cancer. A polyvalent or polyclonal CY cancer, melanoma or epithelial cancer. A polyvalent or polyclonal CY cancer, infectious diseases and promoting beneficial autoimmune CY reating cancer, infectious diseases and promoting beneficial autoimmune CY components. The polyvalent or polyclonal preparations are useful for treating autoimmune diseases with inflammatory or allergic components. The polyvalent or polyclonal preparations are useful for treating autoimmune diseases such as rheumatoid arthritis myssthenia CY diabetes mellitus, transplant rejection, and inflammation, graft versus
                                  Query Match
Best Local Similarity
Matches 9; Conser
                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 26; Fig 8; 127pp; English
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lymphocyte associated antigen-4, useful for inducing, augmenting
prolonging immune response to antigen or for suppressing immune
                                                                                                                                            host disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MEDA-) MEDAREX INC.
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 TGWLGPFDY 9
                                                                                                          118
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                                     Conservative
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                                                  100.0%; Score 58; 100.0%; Pred. No.
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                                    Mismatches
                                                    0.014;
                                                                    DB 22;
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                                                                  Length 118;
                                  Indels
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Novel antibodies capable of binding (CTLA)-4 containing specified heavy for treating, e.g. immune disorders

cytotoxic T-lymphocyte antigen and light chain sequences, use

useful

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RESULT 8
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WPI; 2000-442647/38
N-PSDB; AAA46888.
                                                                                                                                                                                                                                                                                                                                                                                        Cytotoxic T-lymphocyte antigen-4; CTLA-4; antibody; immune system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY93725;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY93725 standard; Protein; 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               a cluster of methionine and charged residues at its C-terminal side. This polymer exhibits an amphiphillic structure and is used as a stabiliser for oil-in-water emulsions. The oil-in-water emulsion is used for producing foodstuffs, pharmaceutical products or cosmetic products by combining with nutritionally, pharmaceutically, and cosmetically suitable ingredients. It is also useful in photography.
                                                        Corvalan JR;
                                                                            Hanson DC,
                                                                                                                                                                   23-DEC-1998;
                                                                                                                                                                                                      23-DEC-1999;
                                                                                                                                                                                                                                               29-JUN-2000
                                                                                                                                                                                                                                                                                  WO200037504-A2
                                                                                                                                                                                                                                                                                                                                                          proliferative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is a synthetic amphiphilic recombinant collagen-
like (or gelatin-like) polymer, N1N2P4 which contains two different
yn-polar modules (N1) and four polar modules (P4). The N2 module
s similar to the N1 module, but differs mainly in the presence of
                                                                                                             (PFIZ ) PFIZER INC. (ABGE-) ABGENIX INC
                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                            hyperimmunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oil-in-water emulsions for preparing foodstuffs, pharmaceutical product or cosmetic product, comprises recombinant collagen-like polymer as stabilizer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Olijve JH, Bo
Wind RD, Van
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GWLGPFDY 9
||||||
142 GWLGPFGY 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 24-26; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-125578/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FUJF ) FUJI PHOTO FILM BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-JUN-2000; 2000EP-0202206
                                                                                                                                                                                                                                                                                                                                                                                                                             ue heavy chain of immunoglobulin clone 12.9.1.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 7; Conser
                                                                        Neveu MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             599 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bouwstra JB, De
an Den Bosch TJ,
                                                                                                                                                                                                                                                                                                                                                          disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                   98US-0113647
                                                                                                                                                                                                        99WO-US30895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99EP-0202047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       79.3%;
87.5%;
                                                                          Mueller
                                                                                                                                                                                                                                                                                                                                                        autoimmune cancer; imm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wolf FA,
Toda Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 46; I
Pred. No. 9
                                                                        E
                                                                                                                                                                                                                                                                                                                                                       immunodeficient disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                          disease;
                                                                        Hanke.
                                                                                                                                                                                                                        `
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22; Length 599;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Werten MWT,
                                                                      Ë
                                                                                                                                                                                                                                                                                                                                                                          diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -
                                                                        Gilman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wisselink HW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in photography
                                                                    SC,
                                                                                                                                                                                                                                                                                                                                                                        graft
                                                                    Davis CG
                                                                                                                                                                                                                                                                                                                                                                        rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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밁 Ś

The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived for the method involves contacting

ID NO 11580; 327pp + sequence listing; English

New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -

Claim

27;

SEQ

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RESULT 9
ABB28612
ID ABB2
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                  04-FEB-2000; 2000US-0180312.

26-MAY-2000; 2000US-0207456.

30-JUN-2000; 2000US-0608408.

03-AUG-2000; 2000US-0632368.

21-SEP-2000; 2000US-0234687.

27-SEP-2000; 2000US-0236359.

04-OCT-2000; 2000GB-0024263.
                                                                                              WPI; 2001-496933/54.
                                                                                                                                                                                                                                                                                                                                                                                                   Peptide #1263 encoded by breast cell single exon nucleic acid probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CTLA) 4. Antibodies of the invention are composed of a heavy chain variable region, comprising a modified contiguous sequence from a FR1-FR3 sequence encoded by a human VH3-33 family gene. The modifications are contained in CDR1, CDR2 and/or framework regions. The antibodies may be used to inhibit CTLA-4 and down-regulate the immune system to treat hyperimmunity disorders (e.g. autoimmune disease diabetes and graft rejection) and proliferative disorders (e.g. cancer). CTLA-4 stimulatory agents may be used to up-regulate immune system to up-regulate immunodeficient disorders.
                                                                                                                       Penn
                                                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US00662
                                                                                                                                                                                                                                                                                              09-AUG-2001.
                                                                                                                                                                                                                                                                                                                       WO200157271-A2
                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                      disease; cancer.
                                                                                                                                               (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                              Human; microarray; single exon probe; gene expression; breast;
                                                                                                                                                                                                                                                                                                                                                                                                                                   01-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB28612;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABB28612 standard; Peptide; 41 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a heavy chain of an antibody of the invention. The antibody is directed cytotoxic T-lymphocyte antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TGWLGPFDY 9
                                                                                                                     SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGWYGGFDF .99
                                                                                                                                            MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 66. 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fig 1M; 157pp;
                                                                                                                    Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               163 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69.0%;
                                                                                                                    Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 40; DB
Pred. No. 26;
1; Mismatches
                                                                                                                     Rank
                                                                                                                     R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 163;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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RESULT 10
ARB33795
ID ARB33795
AC ARB33
AC ARB33795
AC ARB33
AC ARB33
AC ARB33795
AC ARB33
AC ARB33795
AC ARB33
AC ARB33
AC ARB33795
AC ARB33
AC AR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Macches
     The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a peptide encoded by a single exor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4-FEB-2000; 2000US-0180312.
26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
                                                                                                                                                                                                                                     Human genome-derived single exon nucleic analyzing gene expression in human fetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide encoded by a single exon nucleic acid byto of the invention.
                                                                                                                                                                                       Claim 27; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                  Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001; 2001WO-US00669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200157277-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide #1301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The sequence data for this patent did not form p
printed specification, but was obtained in electronic
from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB33795 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     y Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGWLGP---FDY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGWLGPVSIFGY 12
                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                               Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoded by human foetal liver single exon probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           liver;
                                                                                                                                                                                     NO 26430;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene expression; single exon nucleic acid probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68.1%;
                                                                                                                                                                                                                                                                                                                                                                             Chen
                                                                                                                                                                            639pp + sequence listing; English
                                                                                                                                                                                                                                                                                                                                                                          Ę
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
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Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                             Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                          DR
                                                                                                                                                                                                                                        acid probes useful
liver -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               part of the
c format directly
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                                                                                                                                                                                                                                                              for
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acid

probe

Query Match Best Local Similarity

68.1**%**; 66.7**%**;

Score 39.5; D Pred. No. 7.5;

DB: 22;

Length

41;

Sequence

41

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RESULT 11
ABB19238
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                      hypertension, cardiac arrhythmias and congenital heart disease Note: The sequence data for this patent did not form part of the specification, but was obtained in electronic format directly the feature of the feature
                                                                                                                               By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of human heart and vascular system e.g. cardiovascular disease,
                                                                                                                                                                                                ABA21535-ABA41305). The present sequence is a protein encoded by one sucprobe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays
                                                                                                                                                                                                                                                                           The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart
                                                                                                                                                                                                                                                                                                                                                               Claim
                                          at ftp.wipo.int/pub/published
                                                                                                                                                                                                                                                                                                                                                                                                                                   Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-488899/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Penn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001; 2001WO-US00666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; gene expression; heart; microarray; cardiovascular disease; hypertension; cardi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200157274-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  congenital
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB19238 standard; Protein; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB19238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MOLE-)
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; 2000US-0207456.
; 2000US-0608408.
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- 6/ptodata/1/pubpaa/US09 NEW PUB.pep:*
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US-09-801-574-12
US-09-740-668A-14
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paa/US07_NEW_PUB.pep:*
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Sequence 3877, Ap
Sequence 2958, App
Sequence 70, Appl
Sequence 1526, Ap
Sequence 1510, Ap
Sequence 1519, Ap
Sequence 6656, Ap
Sequence 6656, Ap
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Sequence 23, App
Sequence 12, Appl
Sequence 14, Appl
Sequence 70, Appl
Sequence 10380, A
Sequence 11155, A
Sequence 13798, A
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Sequence 36870, A
Sequence 15
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GENERAL INFORMATION:

APPLICANT: XU, Y1

APPLICANT: HOOK, Magnus A.O.

TITLE OF INVENTION: SURFACE PROTEINS FROM GRAM-POSITIVE BACTERIA

FILE REFERENCE: P07196US01/BAS

CURRENT APPLICATION NUMBER: US/10/140,372

CURRENT FILING DATE: 2002-05-08

PRIOR APPLICATION NUMBER: US 60/289,132

PRIOR FILING DATE: 2001-05-08

Sequence 3, Application US/10140372 Publication No. US20030021789A1

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RESULT 2
US-09-864-761-34536
; Sequence 34536, Application US/09864761
patent No. US20020048763A1
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Best Local Similarity 66.,
Thes 6; Conservative.
                                                               APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
                                                                                                                                                                                                                                 239 SAWLNPFDY 247
                                                                                                                                                                                                                                                    1 TGWLGPFDY 9
T: Hanzel, David K.
T: Chen, Wensheng
INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
ERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                72.4%;
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Pred. No. 23
                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                               Length 408;
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; ORGANISM: Streptococcus mutans US-10-140-372-3

SEQ ID NO 3

LENGTH: 408 TYPE: PRT NUMBER OF SEQ ID NOS: 16 SOFTWARE: PatentIn version

version 3.1

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US-09-864-761-36870
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Best Local Similarity
Matches 8; Conserv
Sequence 36870, Application US/09864761 Patent No. US20020048763A1
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EQ ID NO 34536
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens FEATURE:
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FRENT FILING DATE: 2001-05-23
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CATION NUMBER: US 60/236,359
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PATION NUMBER: US 09/608,408
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Pred. No. 6.9;
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ADULT LIVER, SIGNAL =
T: AU137038.1, EVALUE 1
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EVALUE 1.00e-18
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LENGTH: 67
                                                                       Matches
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Best Local (
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PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
                                                                                                                                                                                                                             OTHER INFORMATION: MAP TO AL110502. OTHER INFORMATION: EXPRESSED IN HEL
                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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TLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
LE REFERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 2000-06-30
APPLICATION NUMBER: US 09/774,203
FILING DATE: 2001-01-29
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FILING DATE: 2000-05-26
APPLICATION NUMBER: US 09/632,366
FILING DATE: 2000-08-03
                                   GMLGPFDY 9
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APPLICATION NUMBER: US 09/608,408
GWRGAFDY 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/234,687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
                                                                                                                                                                             INFORMATION:
                                                                                                                                                          INFORMATION:
                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PLICATION NUMBER: PCT/US01/0066:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LING DATE: 2001-01-30
PLICATION NUMBER: PCT/US01/00664
LING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PLICATION NUMBER: PCT/US01/00666
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LING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PLICATION NUMBER: PCT/US01/00668
                                                                    Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATION NUMBER: PCT/US01/00662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CATION NUMBER: PCT/US01/00665
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                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ON NUMBER: PCT/US01/00670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ON NUMBER: PCT/US01/00663
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                                                                                                                                                       EST_HUMAN HIT: BE257700.1,
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                                                                                  67.2%;
75.0%;
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IN BT474,
                                                                                  Score 39;
Pred. No.
                                                                Mismatches
                                                                                                                                                                                                                           SIGNAL = 0.93
                                                                                                                                                                     SIGNAL = 1.2
SIGNAL = 1.1
441, EVALUE 3.
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13;
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                                                                                               Length 67;
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RESULT 4
US-09-872-523-15
; Sequence 15, Application US/09872523

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Result
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Perfect score:
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seq length: 2000000000
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Gapop 10.0 , Gapext 0.5
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58
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                                                                US-08-852-153-4
US-08-852-153-8
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US-08-956-597-3
US-08-2597-451-5
US-08-2597-451-5
US-08-259-4910-13
US-07-672-483-4
US-08-259-451-3
US-07-872-20
US-08-383-753-20
US-08-958-772-20
US-08-958-772-20
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US-09-5137-4
US-09-137-422A-32
US-09-134-001C-3112
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Sequence 3, Appli
Sequence 5, Appli
Sequence 8, Appli
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Sequence 13, Appli
Sequence 4, Appli
Sequence 20, Appli
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	3, Appli	11, Appl	16, Appl	2, Appli	16, Appl	 Appli 	3, Appli	3, Appli	•	13, Appl	2, Appli	23, App]	88, App]	15, App]	11, Appl	150, App	144, Api	10, App.

RESULT 1 US-08-852-153-4

ALIGNMENTS

US-08-852-153-4 Query Match Best Local Similarity Matches 6; Conserv Sequence 4, Application US/08852153 Patent No. 5914266 GENERAL INFORMATION: TELEFAX: (510) 655-35 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk TOPOLOGY: 11 SEQUENCE CHARACTERISTICS: LENGTH: 577 amino acids ATTORNEY/AGENT INFORMATION: NAME: Guth, Joseph H. REGISTRATION NUMBER: 31,261 REFERENCE/DOCKET NUMBER: 12 TELECOMMUNICATION INFORMATION: SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: CORRESPONDENCE ADDRESS: ADDRESSEE: Chiron Corporation COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS STRANDEDNESS: APPLICATION NUMBER: FILING DATE: 2 GWLGPFDY 9 TELEPHONE: CLASSIFICATION: 435 T: 4560 Horton Street Emeryville : California RY: U.S.A. 94608 amino acid (510) 923-38 (510) 655-3542 Randazzo, protein single 923-3888 67.2**%**; 75.0**%**; , Filippo Mammalian Sex Comb on Midleg Acts as a Tumor Suppressor 8 US/08/852,153 1224.006 Version #1.30 <u>ب</u> 2; Indels Length 577; 0

RESULT 2

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142 GWRGAFDY 149

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US-08-852-153-6
; Sequence 6, Application US/08852153
; Patent No. 5914266
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURPENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 591 amino acids
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                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                        TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 31,261
REFERENCE/DOCKET NUMBER: 12
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 923-3888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                         156 GWRGAFDY 163
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 URRENT APPLICATION DATA:
                                                  APSTICATION NUMBER:
                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 43:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ORNEY) AGENT INFORMATION:
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                                                                                                                                                                                                                 DDRESSEE: Chiron Corporation [REET: 4560 Horton Street [TY: Emeryville]
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                                                                                                                                                                                                                                                                                                                                                                    Application US/08852153
                                                                                                                                                                                                   California
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                      ATION: 435
                                                                                                                                                                                  U.S.A.
                                                                                                                                                                                                                                                                                                                     Randazzo,
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PENTION: Mammalian Sex Comb on Midleg Acts as a Tumor Suppressor
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75.0%;
                                                                                                                                                                                                                                                                                                                 Filippo
                                                US/08/852;153
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Pred. No. 62;
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Query Match
Best Local Similarity
Thes 6; Conserv:
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Patent No. 5914266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 75. Matches 6; Conservative
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                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                        MOLECULE TYPE:
                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron Corporation
                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                          TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                       COPOLOGY:
                                                                                                                                                                                              TELEPHONE:
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                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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GWLGPFDY 9
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                         Conservative
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/ENTION: Mammalian Sex Comb on Midleg Acts as
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                                    Score 39; DE
Pred. No. 69;
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Pred. No. 65;
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                                                                                                                                                                                                                     224.006
                                                                                                                                                                                                                                                                                                                          Version #1.30
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                                              Length 664;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 620
                     Indels
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                   0;
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                   Gaps
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